18



SEQUENCE LISTING

<110> Ruvkun, Gary.
Ogg, Scott

<120> THERAPEUTIC AND DIAGNOSTIC TOOLS FOR IMPAIRED GLUCOSE TOLERANCE CONDITIONS

- <130> 00786/351004
- <140> 09/205,658
- <141> 1998-12-03
- <150> 08/857,076
- <151> 1997-05-15
- <150> 08/888,534
- <151> 1997-07-07
- <150> US98/10080
- <151> 1998-05-15
- <160> 328
- <170> FastSEQ for Windows Version 4.0
- <210> 1
- <211> 20
- <212> DNA
- <213> Artificial Sequence
- <220>
- <223> Primer/probe derived from C. elegans
- <400> 1
- cgctacggca aaaaagtgaa
- <210> 2
- <211> 18
- <212> DNA
- <213> Artificial Sequence
- <220>
- <223> Primer/probe derived from C. elegans
- <400> 2
- cgatgatgaa gatacccc
- <210> 3
- <211> 20
- <212> DNA
- <213> Artificial Sequence
- <220>
- <223> Primer/probe derived from C. elegans

amdta

```
. <211> 20
 <212> DNA
 <213> Artificial Sequence
 <223> Primer/probe derived from C. elegans
 <400> 9
                                                                          20
 cgctacggca aaaaagtgaa
 <21.0> 10
 <211> 20
 <212> DNA
 <213> Artificial Sequence
 <220>
 <223> Primer/probe derived from C. elegans
 <400> 10
                                                                          20
 gacgatcccg aggtgagtat
 <210> 11
 <211> 5816
 <212> DNA
 <213> Caenorhabditis elegans
 <220>
 <221> misc_feature
 <222> (1) ... (5816)
 <223> n = A, T, C or G
 <400> 11
 ggtttaatta cccaagtttg agctccaaga gcacacatct gatcgtcgga ttctactgta
                                                                          60
 ctccccgaaa aaccaacaaa aaacacaagt ttttgaacac ttgtaaatgc agacagaacg
                                                                         120
 atgacgagaa tgaatattgt cagatgtcgg agacgacaca aaattttgga aaatttggaa
                                                                         180
                                                                         240
 gaagagaatc teggeeegag etgetegteg acgaetteaa caacegetge caeegaaget
 ctcggaacaa ccactgagga tatgaggctt aagcagcagc gaagctcgtc gcgtgccacg
                                                                         300
 gagcacgata ttgtcgacgg caatcaccac gacgacgagc acatcacaat gagacggctt
                                                                         360
 cgacttgtca aaaattcgcg gacgcggcgt agaacgacgc ccgattcaag tatggactgc
                                                                         420
 tatgaggaaa acccgccatc acaaaaactt caataaatta ttcttggatt tctaaaaagt
                                                                         480
 catcaatgac gtcattaatg cttttactgc tattcgcttt tgtacagccg tgtgcctcaa
                                                                         540
 tagtogaaaa acgatgoggo ocaatogata ttogaaatag googtgggat attaagoogo
                                                                         600
 aatggtcgaa acttggtgat ccgaacgaaa aagatttggc tggtcagaga atggtcaact
                                                                         660
                                                                         720
 gcacagtggt ggaaggttcg ctgacaatct catttgtact gaaacacaag acaaaagcac
 aagaagaaat gcatcgaagt ctacagccaa gatattccca agacgaattt atcacttttc
                                                                         780
 cgcatctacg tgaaattact ggaactctgc tcgtttttga gactgaagga ttagtggatt
                                                                         840
 tgcgtaaaat tttcccaaat cttcgtgtaa ttggaggccg ttcgctgatt caacactatg
                                                                         900
 cgctgataat ttatcgaaat ccggatttgg agatcggtct tgacaagctt tccgtaattc
                                                                         960
 gaaatggtgg tgtacggata atcgataatc gaaaactgtg ctacacgaaa acgattgatt
                                                                        1020
 ggaaacattt qatcacttct tccatcaacq atqttqtcqt tqataatqct qccqaqtacq
                                                                        1080
 ctgtcactga gactggattg atgtgcccac gtggagcttg cgaagaggat aaaggcgaat
                                                                        1140
 caaagtgtca ttatttggag gaaaagaatc aggaacaagg tgtcgaaaga gttcagagtt
                                                                        1200
 gttggtcgaa caccacttgc caaaagtctt gtgcttatga tcgtcttctt ccaacgaaag
                                                                        1260
 adatoggaco gggatgtgat gogaacggog atogatgtoa ogatoaatgo gtgggoggtt
                                                                        1320
 gtgagcgtgt gaatgatgcc acagcatgcc acgcgtgcaa gaatgtctat cacaagggaa
                                                                        1380
 agtgtatcga aaagtgtgat gctcacctgt accttctcct tcaacgtcgt tgtgtgaccc
                                                                        1440
 gtgagcagtg tctgcagctg aatccggtgc tctcgaacaa aacagtgcct atcaaggcga
                                                                        1500
 eggeaggeet ttgeteggat aaatgteeeg atggttatea aateaaceeg gatgateate
                                                                        1560
 gagaatgccg aaaatgcgtt ggcaagtgtg agattgtgtg cgagatcaat cacgtcattg
                                                                        1620
```

atacgtttcc gaaggcacag gcgatcaggc tatgcaatat tattgacgga aatctgacga 1680 tcgagattcg cggaaaacag gattcgggaa tggcgtccga gttgaaggat atatttgcga 1740 acattcacac gatcaccggc tacctgttgg tacgtcaatc gtcaccgttt atctcgttga 1800 1860 acatgttccg gaatttacga cgtattgagg caaagtcact gttcagaaat ctatatgcta 1920 tcacagtttt tgaaaatccg aatttaaaaa agctattcga ttcaacgacg gatttgacgc ttgatcgtgg aactgtgtca attgccaata acaagatgtt atgcttcaag tatatcaagc 1980 agctaatgtc aaagttaaat ataccactcg atccgataga tcaatcagaa gggacaaatg 2040 gtgagaaggn aatctgtgag gatatggcaa tcaacgtgag catcacagcg gtcaacgcgg 2100 actoggtott otttagttgg cootcattca acattacoga tatagatcag ogaaaqttto 2160 2220 teggetaega getettette aaagaagtee caegaatega tgagaacatg aegategaag aggatcgaag tqcgtqtqtc qattcgtqqc agaqtqtctt caaacaqtac tacqaqacqt 2280 cgaacggtga accgaccccg gacattttta tggatattgg accgcgcgag cgaattcggc 2340 cgaatacgct ctacgcgtac tatgtggcga cgcagatggt gttgcatgcc ggtgcgaaga 2400 acggtqtatc qaaqattqqt tttqtqaqqa cqaqctacta tacqcctqat cctccqacqt 2460 tggcactagc gcaagtcgat tcggacgcta ttcatattac gtgggaagcg ccgctccaac 2520 cgaacggaga cctcacgcat tacacaatta tgtggcgtga gaatgaagtg agcccgtacg 2580 aggaagccga aaagttttgt acagatgcaa gcaccccgc aaatcgacaa cgcacgaaag 2640 atccgaaaga gacgattgta gccgataagc cagtcgatat tccgtcatca cgtaccgtag 2700 ctccgacact tttgactatg atgggtcacg aagatcagca gaaaacgtgc gctgcaacgc 2760 ccggttgttg ttcgtgttcg gctatcgaag aatcatcgga acagaacaag aagaagcgac 2820 cggatccgat gtcggcgatc gaatcatctg catttgagaa taagctgttg gatgaggttt 2880 taatgccgag agacacgatg cgagtgagac gatcaattga agacgcgaat cgagtcagtg 2940 aagagttgga aaaagctgaa aatttgggaa aagctccaaa aactctcggt ggaaagaagc 3000 cgctgatcca tatttcgaag aagaagccgt cgagcagcag caccacatcc acaccggctc 3060 caacgatcgc atcaatgtat gccttaacaa ggaaaccgac tacggtgccg ggaacaagga 3120 ttcggctcta cgagatctac gaacctttac ccggaagctg ggcgattaat gtatcagctc 3180 tggcattgga taatagttat gtgatacgaa atttgaagca ttacacactt tatgcgattt 3240 ctctatccgc gtgccaaaac atgacagtac ccggagcatc ttgctcaata tcccatcgtg 3300 cgggagcatt gaaacgaaca aaacacatca cagacattga taaagtgttg aatgaaacaa 3360 ttgaatggag atttatgaat aatagtcaac aagtcaacgt gacgtgggat ccaccgactg 3420 aagtgaatgg tggaatattc ggttatgttg taaagcttaa gtcaaaagtc gatggatcaa 3480 ttgttatgac gagatgtgtc ggtgcgaaga gaggatattc aacacggaat cagggtgtcc 3540 tattccagaa tttggccgat ggacgttatt ttgtctcagt aacggcgacc tctgtacacg 3600 gegetggace ggaageegaa teeteegaee caategtegt catgaegeea ggettettea 3660 ctgtggaaat cattctcggc atgcttctcg tctttttgat tttaatgtca attgccggtt 3720 gtataatcta ctactacatt caagtacgct acggcaaaaa agtgaaagct ctatctgact 3780 ttatgcaatt gaatcccgaa tattgtgtgg acaataagta caatgcagac gattgggagc 3840 tacggcagga tgatgttgtg ctcggacaac agtgtggaga gggatcattc ggaaaagtgt 3900 acctaggaac tggaaataat gttgtttctc tgatgggtga tcgtttcgga ccgtqtgcta 3960 ttaagattaa tgtagatgat ccagcgtcga ctgagaatct caactatctc atggaagcta 4020 atattatgaa qaactttaaq actaacttta tcqtccaact qtacqqaqtt atctctactq 4080 tacaaccagc gatggttgtg atggaaatga tggatcttgg aaatctccgt gactatctcc 4140 4200 gatcgaaacg cgaagacgaa gtgttcaatg agacggactg caactttttc gacataatcc cgagggataa attccatgag tgggccgcac agatttgtga tggtatggcg tacctggagt 4260 4320 cgctcaagtt ttgccatcga gatctcgccg cacgtaattg catgataaat cgggatgaga ctgtcaagat tggagatttc ggaatggctc gtgatctatt ctatcatgac tattataagc 4380 categggcaa gegtatgatg cetgttegat ggatgteace egagtegttg aaagaeggaa 4440 agtttgactc gaaatctgat gtttggagct tcggagttgt tctctatgaa atggttacac 4500 4560 teggtgetea gecatatatt ggtttgagta atgatgaggt gttgaattat attggaatgg 4620 cccggaaggt tatcaagaag cccgaatgtt gtgaaaacta ttggtataag gtgatgaaaa 4680 tgtgctggag atactcacct cgggatcgtc cgacgttcct ccagctcgtt catcttctag cagctgaagc ttcaccagaa ttccgagatt tatcatttgt cctaaccgat aatcaaatga 4740 4800 tccttgacga ttcagaagca ctggatcttg atgatattga tgatactgat atgaatgatc 4860 aggttgtcga ggtggcaccg gatgttgaga acgtcgaggt tcagagtgat tcggaacgtc ggaatacgga ttcaataccg ttgaaacagt ttaagacgat ccctccgatc aatgcgacga 4920 4980 cgagtcattc gacaatatcg attgatgaga caccgatgaa agcgaagcag cgagaaggat cgctggatga ggagtacgca ttgatgaatc atagtggagg tccgagtgat gcggaagttc 5040 ggacgtatgc tggtgatgga gattatgtgg agagagatgt tcgagagaat gatgtgccaa 5100 5160 cgcgacgaaa tactggtgca tcaacatcaa gttacacagg tggtggtcca tattgcctaa

```
caaatcgtgg tggttcaaat gaacgaggag ccggtttcgg tgaagcagta cgattaactq
                                                                      5220
atggtgttgg aagtggacat ttaaatgatg atgattatgt tgaaaaagag atatcatcca
                                                                      5280
tggatacgcg ccggagcacg ggcgcctcga gctcttccta cggtgttcca cagacgaatt
                                                                      5340
ggagtggaaa tcgtggtgcc acgtattata cgagtaaagc tcaacaggca gcaactgcag
                                                                      5400
cagcagcagc agcagcagct ctccaacagc aacaaaatgg tggtcgaggc gatcgattaa
                                                                      5460
ctcaactacc cggaactgga catttacaat cgacacgtgg tggacaagat ggagattata
                                                                      5520
ttgaaactga accgaaaaat tatagaaata atggatctcc atcgcgaaac ggcaacagcc
                                                                      5580
gtgacatttt caacggacgt tcggctttcg gtgaaaatga gcatctaatc gaggataatg
                                                                      5640
agcatcatcc acttgtctga aacccccaaa aaatcccgcc tcttaaatta taaattatct
                                                                      5700
cccacattat catatctcta cacgaatatc ggattttttt tcagattttt tctgaaaaat
                                                                      5760
totgaataat titaccccat tittcaaato totgtatiit tittigitat tacccc
                                                                      5816
```

```
<210> 12
<211> 1724
<212> PRT
<213> Caenorhabditis elegans
```

<400> 12 Met Thr Ser Leu Met Leu Leu Leu Phe Ala Phe Val Gln Pro Cys 1 Ala Ser Ile Val Glu Lys Arg Cys Gly Pro Ile Asp Ile Arg Asn Arg Pro Trp Asp Ile Lys Pro Gln Trp Ser Lys Leu Gly Asp Pro Asn Glu Lys Asp Leu Ala Gly Gln Arg Met Val Asn Cys Thr Val Val Glu Gly Ser Leu Thr Ile Ser Phe Val Leu Lys His Lys Thr Lys Ala Gln Glu Glu Met His Arg Ser Leu Gln Pro Arg Tyr Ser Gln Asp Glu Phe Ile 90 Thr Phe Pro His Leu Arg Glu Ile Thr Gly Thr Leu Leu Val Phe Glu 105 Thr Glu Gly Leu Val Asp Leu Arg Lys Ile Phe Pro Asn Leu Arg Val 120 Ile Gly Gly Arg Ser Leu Ile Gln His Tyr Ala Leu Ile Ile Tyr Arg Asn Pro Asp Leu Glu Ile Gly Leu Asp Lys Leu Ser Val Ile Arg Asn 155 160 Gly Gly Val Arg Ile Ile Asp Asn Arg Lys Leu Cys Tyr Thr Lys Thr 170 Ile Asp Trp Lys His Leu Ile Thr Ser Ser Ile Asn Asp Val Val Val 180 185 190 Asp Asn Ala Ala Glu Tyr Ala Val Thr Glu Thr Gly Leu Met Cys Pro 200 Arg Gly Ala Cys Glu Glu Asp Lys Gly Glu Ser Lys Cys His Tyr Leu 215 Glu Glu Lys Asn Gln Glu Gln Gly Val Glu Arg Val Gln Ser Cys Trp 230 235 Ser Asn Thr Thr Cys Gln Lys Ser Cys Ala Tyr Asp Arg Leu Leu Pro 250 Thr Lys Glu Ile Gly Pro Gly Cys Asp Ala Asn Gly Asp Arg Cys His 265 270 Asp Gln Cys Val Gly Gly Cys Glu Arg Val Asn Asp Ala Thr Ala Cys 275 280 His Ala Cys Lys Asn Val Tyr His Lys Gly Lys Cys Ile Glu Lys Cys 295 300 Asp Ala His Leu Tyr Leu Leu Gln Arg Arg Cys Val Thr Arg Glu

```
315
                     310
 Gln Cys Leu Gln Leu Asn. Pro Val Leu Ser Asn Lys Thr Val Pro Ile
                 325
                                     330
Lys Ala Thr Ala Gly Leu Cys Ser Asp Lys Cys Pro Asp Gly Tyr Gln
          . 340
                                 345
                                                      350
 Ile Asn Pro Asp Asp His Arg Glu Cys Arg Lys Cys Val Gly Lys Cys
                             360
 Glu Ile Val Cys Glu Ile Asn His Val Ile Asp Thr Phe Pro Lys Ala
                         375
                                              380
 Gln Ala Ile Arg Leu Cys Asn Ile Ile Asp Gly Asn Leu Thr Ile Glu
                     390
                                         395
 Ile Arg Gly Lys Gln Asp Ser Gly Met Ala Ser Glu Leu Lys Asp Ile
                 405
                                      410
 Phe Ala Asn Ile His Thr Ile Thr Gly Tyr Leu Leu Val Arg Gln Ser
                                 425
             420
 Ser Pro Phe Ile Ser Leu Asn Met Phe Arg Asn Leu Arg Arg Ile Glu
         435
                             440
 Ala Lys Ser Leu Phe Arg Asn Leu Tyr Ala Ile Thr Val Phe Glu Asn
     450
                         455
 Pro Asn Leu Lys Lys Leu Phe Asp Ser Thr Thr Asp Leu Thr Leu Asp
                     470
                                         475
 Arg Gly Thr Val Ser Ile Ala Asn Asn Lys Met Leu Cys Phe Lys Tyr
                 485
                                      490
 Ile Lys Gln Leu Met Ser Lys Leu Asn Ile Pro Leu Asp Pro Ile Asp
             500
                                 505
 Gln Ser Glu Gly Thr Asn Gly Glu Lys Ala Ile Cys Glu Asp Met Ala
         515
                             520
 Ile Asn Val Ser Ile Thr Ala Val Asn Ala Asp Ser Val Phe Phe Ser
                         535
                                              540
 Trp Pro Ser Phe Asn Ile Thr Asp Ile Asp Gln Arg Lys Phe Leu Gly
                                         555
                     550
 Tyr Glu Leu Phe Phe Lys Glu Val Pro Arg Ile Asp Glu Asn Met Thr
                 565
                                      570
 Ile Glu Glu Asp Arg Ser Ala Cys Val Asp Ser Trp Gln Ser Val Phe
             580
                                 585
 Lys Gln Tyr Tyr Glu Thr Ser Asn Gly Glu Pro Thr Pro Asp Ile Phe
         595
                             600
Met Asp Ile Gly Pro Arg Glu Arg Ile Arg Pro Asn Thr Leu Tyr Ala
     610
                         615
                                              620
 Tyr Tyr Val Ala Thr Gln Met Val Leu His Ala Gly Ala Lys Asn Gly
                     630
                                         635
 Val Ser Lys Ile Gly Phe Val Arg Thr Ser Tyr Tyr Thr Pro Asp Pro
                 645
                                     650
Pro Thr Leu Ala Leu Ala Gln Val Asp Ser Asp Ala Ile His Ile Thr
             660
                                 665
 Trp Glu Ala Pro Leu Gln Pro Asn Gly Asp Leu Thr His Tyr Thr Ile
         675
                             680
Met Trp Arg Glu Asn Glu Val Ser Pro Tyr Glu Glu Ala Glu Lys Phe
                         695
Cys Thr Asp Ala Ser Thr Pro Ala Asn Arg Gln Arg Thr Lys Asp Pro
                     710
                                         715
 Lys Glu Thr Ile Val Ala Asp Lys Pro Val Asp Ile Pro Ser Ser Arg
                 725
                                     730
 Thr Val Ala Pro Thr Leu Leu Thr Met Met Gly His Glu Asp Gln Gln
             740
                                 745
 Lys Thr Cys Ala Ala Thr Pro Gly Cys Cys Ser Cys Ser Ala Ile Glu
                             760
Glu Ser Ser Glu Gln Asn Lys Lys Lys Arg Pro Asp Pro Met Ser Ala
     770
                         775
                                              780
```

```
Ile Glu Ser Ser Ala Phe Glu Asn Lys Leu Leu Asp Glu Val Leu Met
                    790
                                        795
Pro Arg Asp Thr Met Arg Val Arg Arg Ser Ile Glu Asp Ala Asn Arg
                805
                                    810
Val Ser Glu Glu Leu Glu Lys Ala Glu Asn Leu Gly Lys Ala Pro Lys
            820
                                825
Thr Leu Gly Gly Lys Lys Pro Leu Ile His Ile Ser Lys Lys Pro
                            840
                                                845
Ser Ser Ser Ser Thr Thr Ser Thr Pro Ala Pro Thr Ile Ala Ser Met
                        855
                                            860
Tyr Ala Leu Thr Arg Lys Pro Thr Thr Val Pro Gly Thr Arg Ile Arg
                   870
                                        875
Leu Tyr Glu Ile Tyr Glu Pro Leu Pro Gly Ser Trp Ala Ile Asn Val
                885
                                    890
Ser Ala Leu Ala Leu Asp Asn Ser Tyr Val Ile Arg Asn Leu Lys His
            900
                                905
                                                    910
Tyr Thr Leu Tyr Ala Ile Ser Leu Ser Ala Cys Gln Asn Met Thr Val
        915
                            920
                                                925
Pro Gly Ala Ser Cys Ser Ile Ser His Arg Ala Gly Ala Leu Lys Arg
   930
                        935
                                            940
Thr Lys His Ile Thr Asp Ile Asp Lys Val Leu Asn Glu Thr Ile Glu
                    950
                                        955
Trp Arg Phe Met Asn Asn Ser Gln Gln Val Asn Val Thr Trp Asp Pro
                965
                                    970
                                                        975
Pro Thr Glu Val Asn Gly Gly Ile Phe Gly Tyr Val Val Lys Leu Lys
            980
                                985
                                                    990
Ser Lys Val Asp Gly Ser Ile Val Met Thr Arg Cys Val Gly Ala Lys
        995
                            1000
                                                1005
Arg Gly Tyr Ser Thr Arg Asn Gln Gly Val Leu Phe Gln Asn Leu Ala
   1010
                       1015
                                            1020
Asp Gly Arg Tyr Phe Val Ser Val Thr Ala Thr Ser Val His Gly Ala
                   1030
                                        1035
Gly Pro Glu Ala Glu Ser Ser Asp Pro Ile Val Val Met Thr Pro Gly
               1045
                                    1050
Phe Phe Thr Val Glu Ile Ile Leu Gly Met Leu Leu Val Phe Leu Ile
           1060
                                1065
                                                    1070
Leu Met Ser Ile Ala Gly Cys Ile Ile Tyr Tyr Tyr Ile Gln Val Arg
       1075
                            1080
                                                1085
Tyr Gly Lys Lys Val Lys Ala Leu Ser Asp Phe Met Gln Leu Asn Pro
   1090
                       1095
                                            1100
Glu Tyr Cys Val Asp Asn Lys Tyr Asn Ala Asp Asp Trp Glu Leu Arg
                   1110
                                        1115
Gln Asp Asp Val Val Leu Gly Gln Gln Cys Gly Glu Gly Ser Phe Gly
               1125
                                    1130
Lys Val Tyr Leu Gly Thr Gly Asn Asn Val Val Ser Leu Met Gly Asp
           .1140
                                1145
Arg Phe Gly Pro Cys Ala Ile Lys Ile Asn Val Asp Asp Pro Ala Ser
       1155
                            1160
                                                1165
Thr Glu Asn Leu Asn Tyr Leu Met Glu Ala Asn Ile Met Lys Asn Phe
   1170
                       1175
                                            1180
Lys Thr Asn Phe Ile Val Gln Leu Tyr Gly Val Ile Ser Thr Val Gln
1185
                   1190
                                        1195
Pro Ala Met Val Val Met Glu Met Asp Leu Gly Asn Leu Arg Asp
            . 1205
                                    1210
Tyr Leu Arg Ser Lys Arg Glu Asp Glu Val Phe Asn Glu Thr Asp Cys
           1220
                                1225
Asn Phe Phe Asp Ile Ile Pro Arg Asp Lys Phe His Glu Trp Ala Ala
                            1240
Gln Ile Cys Asp Gly Met Ala Tyr Leu Glu Ser Leu Lys Phe Cys His
```

1250	1255	5	1260
Arg Asp Leu Ala 1265			n Arg Asp Glu Thr Val
Lys Ile Gly Asp	Phe Gly Met 1285	Ala Arg Asp Let 1290	u Phe Tyr His Asp Tyr 1295
Tyr Lys Pro Ser 130		Met Met Pro Val 1305	l Arg Trp Met Ser Pro 1310
Glu Ser Leu Lys 1315	Asp Gly Lys	Phe Asp Ser Lys 1320	s Ser Asp Val Trp Ser 1325
Phe Gly Val Val 1330	Leu Tyr Glu 1335		u Gly Ala Gln Pro Tyr 1340
Ile Gly Leu Ser 1345	Asn Asp Glu 1350	Val Leu Asn Tyr 13	r Ile Gly Met Ala Arg 55 136
Lys Val Ile Lys	Lys Pro Glu 1365	Cys Cys Glu Asr 1370	n Tyr Trp Tyr Lys Val 1375
Met Lys Met Cys 1380		Ser Pro Arg Asp 1385	o Arg Pro Thr Phe Leu 1390
Gln Leu Val His 1395	Leu Leu Ala	Ala Glu Ala Sei 1400	r Pro Glu Phe Arg Asp 1405
Leu Ser Phe Val 1410	Leu Thr Asp 1415		e Leu Asp Asp Ser Glu 1420
Ala Leu Asp Leu 1425	Asp Asp Ile 1430	Asp Asp Thr Asp 143	p Met Asn Asp Gln Val 35 144
Val Glu Val Ala	Pro Asp Val 1445	Glu Asn Val Glu 1450	u Val Gln Ser Asp Ser 1455
Glu Arg Arg Asn 1460		Ile Pro Leu Lys 1465	s Gln Phe Lys Thr Ile 1470
Pro Pro Ile Asn 1475		Ser His Ser Thi 1480	r Ile Ser Ile Asp Glu 1485
Thr Pro Met Lys 1490	Ala Lys Gln 1495		r Leu Asp Glu Glu Tyr 1500
1505	1510	151	
	1525	1530	p Val Arg Glu Asn Asp 1535
1540)	1545	r Ser Ser Tyr Thr Gly 1550
1555	-	1560	y Ser Asn Glu Arg Gly 1565
1570	1575	5	o Gly Val Gly Ser Gly 1580
1585	1590	159	
	1605	1610	r Tyr Gly Val Pro Gln 1615
1620	0	1625	r Tyr Thr Ser Lys Ala 1630
1635		1640	a Ala Ala Leu Gln Gln 1645
1650	1655	,	r Gln Leu Pro Gly Thr 1660
1665	1670	167	
	1685	1690	r Pro Ser Arg Asn Gly 1695
1700)	1705	a Phe Gly Glu Asn Glu 1710
His Leu Ile Glu 1715	_	His His Pro Let 1720	ı Val

```
<210> 13
<211> 139
<212> PRT
<213> Caenorhabditis elegans
<400> 13
Thr Ser Gly Ser Gly Met Gly Pro Thr Thr Leu His Lys Leu Thr Ile
Gly Gly Gln Ile Arg Leu Thr Gly Arg Val Gly Ser Gly Arg Phe Gly
            20
Asn Val Ser Arg Gly Asp Tyr Arg Gly Glu Ala Val Ala Val Lys Val
Phe Asn Ala Leu Asp Glu Pro Ala Phe His Lys Glu Thr Glu Ile Phe
                        55
Glu Thr Arg Met Leu Arg His Pro Asn Val Leu Arg Tyr Ile Gly Ser
                    70
Asp Arg Val Asp Thr Gly Phe Val Thr Glu Leu Trp Leu Val Thr Glu
Tyr His Pro Ser Gly Ser Leu His Asp Phe Leu Leu Glu Asn Thr Val
                                105
Asn Ile Glu Thr Tyr Tyr Asn Leu Met Arg Ser Thr Ala Ser Gly Leu
                            120
Ala Phe Leu His Asn Gln Ile Gly Gly Ser Lys
    130
                        135
<210> 14
<211> 62
<212> PRT
<213> Caenorhabditis elegans
<400> 14
Glu Asp Ala Ala Ser Asp Ile Ile Ala Asn Glu Asn Tyr Lys Cys Gly
1
Thr Val Arg Tyr Leu Ala Pro Glu Ile Leu Asn Ser Thr Met Gln Phe
            20
Thr Val Phe Glu Ser Tyr Gln Cys Ala Asp Val Tyr Ser Phe Ser Leu
Val Met Trp Glu Thr Leu Cys Arg Cys Glu Asp Gly Asp Val
    50
                        55
<210> 15
<211> 31
<212> PRT
<213> Caenorhabditis elegans
<400> 15
Lys Pro Ala Met Ala His Arg Asp Ile Lys Ser Lys Asn Ile Met Val
Lys Asn Asp Leu Thr Cys Ala Ile Gly Asp Leu Gly Leu Ser Leu
            20
<210> 16
<211> 72
```

<212> PRT

<213> Caenorhabditis elegans

```
<400> 16
Ile Pro Tyr Ile Glu Trp Thr Asp Arg Asp Pro Gln Asp Ala Gln Met
                                    10
Phe Asp Val Val Cys Thr Arg Arg Leu Arg Pro Thr Glu Asn Pro Leu
                                25
Trp Lys Asp His Pro Glu Met Lys His Ile Met Glu Ile Ile Lys Thr
                            40
Cys Trp Asn Gly Asn Pro Ser Ala Arg Phe Thr Ser Tyr Ile Cys Arg
                        55
Lys Arg Met Asp Glu Arg Gln Gln
65
<210> 17
<211> 150
<212> PRT
<213> Caenorhabditis elegans
<400> 17
Tyr Phe Glu Ser Val Asp Arg Phe Leu Tyr Ser Cys Val Gly Tyr Ser
1
Val Ala Thr Tyr Ile Met Gly Ile Lys Asp Arg His Ser Asp Asn Leu
            20
                                25
Met Leu Thr Glu Asp Gly Lys Tyr Val His Ile Asp Phe Gly His Ile
                            40
Leu Gly His Gly Lys Thr Lys Leu Gly Ile Gln Arg Asp Arg Gln Pro
    50
                        55
Phe Ile Leu Thr Glu His Phe Met Thr Val Ile Arg Ser Gly Lys Ser
                    70
Val Asp Gly Asn Ser His Glu Leu Gln Lys Phe Lys Thr Leu Cys Val
                85
                                    90
Glu Ala Tyr Glu Val Met Trp Asn Asn Arg Asp Leu Phe Val Ser Leu
            100
                                105
Phe Thr Leu Met Leu Gly Met Glu Leu Pro Glu Leu Ser Thr Lys Ala
                            120
Asp Leu Asp His Leu Lys Lys Thr Leu Phe Cys Asn Gly Glu Ser Lys
                        135
Glu Glu Ala Arg Lys Phe
<210> 18
<211> 113
<212> PRT
<213> Caenorhabditis elegans
<400> 18
Ser Pro Leu Asp Pro Val Tyr Lys Leu Gly Glu Met Ile Ile Asp Lys
Ala Ile Val Leu Gly Ser Ala Lys Arg Pro Leu Met Leu His Trp Lys
                                25
Asn Lys Asn Pro Lys Ser Asp Leu His Leu Pro Phe Cys Ala Met Ile
Phe Lys Asn Gly Asp Asp Leu Arg Gln Asp Met Leu Val Leu Gln Val
Leu Glu Val Met Asp Asn Ile Trp Lys Ala Ala Asn Ile Asp Cys Cys
                    70
```

Leu Asn Pro Tyr Ala Val Leu Pro Met Gly Glu Met Ile Gly Ile Ile ·

```
Glu Val Val Pro Asn Cys Lys Thr Ile Phe Glu Ile Gln Val Gly Thr
                                105
Gly
<210> 19
<211> 106
<212> PRT
<213> Caenorhabditis elegans
<400> 19
Leu Ala Phe Val Trp Thr Asp Arg Glu Asn Phe Ser Glu Leu Tyr Val
                                                         15
                                    10
Met Leu Glu Lys Trp Lys Pro Pro Ser Val Ala Ala Ala Leu Thr Leu
                                25
Leu Gly Lys Arg Cys Thr Asp Arg Val Ile Arg Lys Phe Ala Val Glu
        35
                            40
Lys Leu Asn Glu Gln Leu Ser Pro Val Thr Phe His Leu Phe Ile Leu
                        55
Pro Leu Ile Gln Ala Leu Lys Tyr Glu Pro Arg Ala Gln Ser Glu Val
                                        75
                    70
Gly Met Met Leu Leu Thr Arg Ala Leu Cys Asp Tyr Arg Ile Gly His
               8.5
                                    90
Arg Leu Phe Trp Leu Leu Arg Ala Glu Ile
            100
<210> 20
<211> 139
<212> PRT
<213> Caenorhabditis elegans
<400> 20
Glu Tyr Trp Ile Val Thr Glu Phe His Glu Arg Leu Ser Leu Tyr Glu
                                    10
Leu Leu Lys Asn Asn Val Ile Ser Ile Thr Ser Ala Asn Arg Ile Ile
                                25
Met Ser Met Ile Asp Gly Leu Gln Phe Leu His Asp Asp Arg Pro Tyr
                            40
Phe Phe Gly His Pro Lys Lys Pro Ile Ile His Arg Asp Ile Lys Ser
                        55
Lys Asn Ile Leu Val Lys Ser Asp Met Thr Thr Cys Ile Ala Asp Phe
                    70
                                        75
Gly Leu Ala Arg Ile Tyr Ser Tyr Asp Ile Glu Gln Ser Asp Leu Leu
                85
                                    90
Gly Gln Val Gly Thr Lys Arg Tyr Met Ser Pro Glu Met Leu Glu Gly
            100
                                105
                                                     110
Ala Thr Glu Phe Thr Pro Thr Ala Phe Lys Ala Met Asp Val Tyr Ser
        115
                            120
Met Gly Leu Val Met Trp Glu Val Ile Ser Arg
    130
                        135
```

```
<210> 21
<211> 61
<212> PRT
```

<213> Caenorhabditis elegans

```
<400> 21
Ile Gly Phe Asp Pro Thr Ile Gly Arg Met Arg Asn Tyr Val Val Ser
Lys Lys Glu Arg Pro Gln Trp Arg Asp Glu Ile Ile Lys His Glu Tyr
Met Ser Leu Lys Lys Val Thr Glu Glu Met Trp Asp Pro Glu Ala
Cys Ala Arg Ile Thr Ala Gly Cys Ala Phe Ala Arg Val
                        55
<210> 22
<211> 20
<212> PRT
<213> Caenorhabditis elegans
<400> 22
Pro Ile Thr Asp Phe Gln Leu Ile Ser Lys Gly Arg Phe Gly Lys Val
1
Phe Lys Ala Gln
<210> 23
<211> 163
<212> PRT
<213> Caenorhabditis elegans
<400> 23
Thr Asp Ser Glu Thr Arg Ser Arg Phe Ser Leu Gly Trp Tyr Asn Asn
Pro Asn Arg Ser Pro Gln Thr Ala Glu Val Arg Gly Leu Ile Gly Lys
                                25
Gly Val Arg Phe Tyr Leu Leu Ala Gly Glu Val Tyr Val Glu Asn Leu
                            40
Cys Asn Ile Pro Val Phe Val Gln Ser Ile Gly Ala Asn Met Lys Asn
                        55
Gly Phe Gln Leu Asn Thr Val Ser Lys Leu Pro Pro Thr Gly Thr Met
                                        75
Lys Val Phe Asp Met Arg Leu Phe Ser Lys Gln Leu Arg Thr Ala Ala
                                    90
Glu Lys Thr Tyr Gln Asp Val Tyr Cys Leu Ser Arg Met Cys Thr Val
            100
                                105
                                                    110
Arg Val Ser Phe Cys Lys Gly Trp Gly Glu His Tyr Arg Arg Ser Thr
                                                125
                            120
Val Leu Arg Ser Pro Val Trp Phe Gln Ala His Leu Asn Asn Pro Met
                        135
                                            140
His Trp Val Asp Ser Val Leu Thr Cys Met Gly Ala Pro Pro Arg Ile
                . 150
Cys Ser Ser
<210> 24
<211> 44
<212> PRT
<213> Caenorhabditis elegans
```

<400> 24

Arg Ala Phe Arg Phe Pro Val Ile Arg Tyr Glu Ser Gln Val Lys Ser 1 10 Ile Leu Thr Cys Arg His Ala Phe Asn Ser His Ser Arg Asn Val Cys 20 25 Leu Asn Pro Tyr His Tyr Arg Trp Val Glu Leu Pro 35 40 <210> 25 <211> 38 <212> PRT <213> Caenorhabditis elegans <400> 25 Val Glu Tyr Glu Glu Ser Pro Ser Trp Leu Lys Leu Ile Tyr Tyr Glu 15 Glu Gly Thr Met Ile Gly Glu Lys Ala Asp Val Glu Gly His His Cys Leu Ile Asp Gly Phe Thr 35 <210> 26 <211> 60 <212> PRT <213> Caenorhabditis elegans <400> 26 Asn Leu Ala Glu Thr Gly His Ser Lys Ile Met Arg Ala Ala His Lys Val Ser Asn Pro Glu Ile Gly Tyr Cys Cys His Pro Thr Glu Tyr Asp 25 Tyr Ile Lys Leu Ile Tyr Val Asn Arg Asp Gly Arg Val Ser Ile Ala 35 Asn Val Asn Gly Met Ile Ala Lys Lys Cys Gly Cys <210> 27 <211> 20 <212> PRT <213> Caenorhabditis elegans <400> 27 Asp Trp Ile Val Ala Pro Pro Arg Tyr Asn Ala Tyr Met Cys Arg Gly 1 15 Asp Cys His Tyr 20 <210> 28 <211> 43 <212> PRT <213> Caenorhabditis elegans

<400> 28

Val Cys Asn Ala Glu Ala Gln Ser Lys Gly Cys Cys Leu Tyr Asp Leu 1 5 10 15 Glu Ile Glu Phe Glu Lys Ile Gly Trp Asp Trp Ile Val Ala Pro Pro

```
20
                                                     30
                                 25
Arg Tyr Asn Ala Tyr Met Cys Arg Gly Asp Cys
        35
                             40
<210> 29
<211> 70
<212> PRT
<213> Caenorhabditis elegans
<400> 29
Asp Cys His Tyr Asn Ala His His Phe Asn Leu Ala Glu Thr Gly His
1
Ser Lys Ile Met Arg Ala Ala His Lys Val Ser Asn Pro Glu Ile Gly
                                 25
            20
Tyr Cys Cys His Pro Thr Glu Tyr Asp Tyr Ile Lys Leu Ile Tyr Val
        35
                             40
Asn Arg Asp Gly Arg Val Ser Ile Ala Asn Val Asn Gly Met Ile Ala
    50
Lys Lys Cys Gly Cys Ser
65
<210>.30
<211> 35
<212> PRT
<213> Caenorhabditis elegans
<400> 30
Cys Cys Leu Tyr Asp Leu Glu Ile Glu Phe Glu Lys Ile Gly Trp Asp
1
Trp Ile Val Ala Pro Pro Arg Tyr Asn Ala Tyr Met Cys Arg Gly Asp
            20
                                 25
                                                     30
Cys His Tyr
        35
<210> 31
<211> 23
<212> DNA
<213> Artificial Sequence
<220>
<223> Degenerate probe
<221> misc_feature
<222> (1)...(23)
<223> n = A, T, C or G
<400> 31
ggntgggayt rnrtnrtngc ncc
                                                                         23
<210> 32
<211> 18
<212> DNA
<213> Artificial Sequence
<220>
<223> Degenerate probe
```

```
<221> misc_feature
<222> (1)...(18)
<223> n = A,T,C or G
<400> 32
tgytgynnnc cnacngar
<210> 33
<211> 127
<212> PRT
<213> Caenorhabditis elegans
<400> 33
Lys Phe His Glu Trp Ala Ala Gln Ile Cys Asp Gly Met Ala Tyr Leu
1
Glu Ser Leu Lys Phe Cys His Arg Asp Leu Ala Ala Arg Asn Cys Met
            20
                                25
Ile Asn Arg Asp Glu Thr Val Lys Ile Gly Asp Phe Gly Met Ala Arg
                            40
Asp Leu Phe Tyr His Asp Tyr Tyr Lys Pro Ser Gly Lys Arg Met Met
                        55
Pro Val Arg Trp Met Ser Pro Glu Ser Leu Lys Asp Gly Lys Phe Asp
                    70
                                        75
Ser Lys Ser Asp Val Trp Ser Phe Gly Val Val Leu Tyr Glu Met Val
                85
                                    90
Thr Leu Gly Ala Gln Pro Tyr Ile Gly Leu Ser Asn Asp Glu Val Leu
            100
                                105
Asn Tyr Ile Gly Met Ala Arg Lys Val Ile Lys Lys Pro Glu Cys
        115
                            120
<210> 34
<211> 131
<212> PRT
<213> Caenorhabditis elegans
<400> 34
Asn Thr Thr Cys Gln Lys Ser Cys Ala Tyr Asp Arg Leu Leu Pro Thr
1
Lys Glu Ile Gly Pro Gly Cys Asp Ala Asn Gly Asp Arg Cys His Asp
            20
Gln Cys Val Gly Gly Cys Glu Arg Val Asn Asp Ala Thr Ala Cys His
Ala Cys Lys Asn Val Tyr His Lys Gly Lys Cys Ile Glu Lys Cys Asp
   50
                        55
Ala His Leu Tyr Leu Leu Gln Arg Arg Cys Val Thr Arg Glu Gln
                    70
Cys Leu Gln Leu Asn Pro Val Leu Ser Asn Lys Thr Val Pro Ile Lys
                                    90
Ala Thr Ala Gly Leu Cys Ser Asp Lys Cys Pro Asp Gly Tyr Gln Ile
                                105
Asn Pro Asp Asp His Arg Glu Cys Arg Lys Cys Val Gly Lys Cys Glu
        115
                            120
Ile Val Cys
   130
```

<210> 35 <211> 103

```
<212> PRT
<213> Caenorhabditis elegans
<400> 35
Phe Asp Gln Lys Ala Cys Glu Ser Leu Val Lys Lys Leu Lys Asp Lys
Lys Asn Asp Leu Gln Asn Leu Ile Asp Val Val Leu Ser Lys Gly Thr
Lys Tyr Thr Gly Cys Ile Thr Ile Pro Arg Thr Leu Asp Gly Arg Leu
Gln Val His Gly Arg Lys Gly Phe Pro His Val Val Tyr Gly Lys Leu
Trp Arg Phe Asn Glu Met Thr Lys Asn Glu Thr Arg His Val Asp His
                                        75
Cys Lys His Ala Phe Glu Met Lys Ser Asp Met Val Cys Val Asn Pro
Tyr His Tyr Glu Ile Val Ile
            100
<210> 36
<211> 79
<212> PRT
<213> Caenorhabditis elegans
<400> 36
Asn Arg Tyr Ser Leu Gly Leu Glu Pro Asn Pro Ile Arg Glu Pro Val
Ala Phe Lys Val Arg Lys Ala Ile Val Asp Gly Ile Arg Phe Ser Tyr
                                25
Lys Lys Asp Gly Ser Val Trp Leu Gln Asn Arg Met Lys Tyr Pro Val
                            40
Phe Val Thr Ser Gly Tyr Leu Asp Glu Gln Ser Gly Gly Leu Lys Lys
                        55
Asp Lys Val His Lys Val Tyr Gly Cys Ala Ser Ile Lys Thr Phe
<210> 37
<211> 106
<212> PRT
<213> Caenorhabditis elegans
<400> 37
Lys Lys Thr Thr Arg Arg Asn Ala Trp Gly Asn Met Ser Tyr Ala
                                    10
Glu Leu Ile Thr Thr Ala Ile Met Ala Ser Pro Glu Lys Arg Leu Thr
                                25
Leu Ala Gln Val Tyr Glu Trp Met Val Gln Asn Val Pro Tyr Phe Arg
                            40
                                                 4.5
Asp Lys Gly Asp Ser Asn Ser Ser Ala Gly Trp Lys Asn Ser Ile Arg
                        55
His Asn Leu Ser Leu His Ser Arg Phe Met Arg Ile Gln Asn Glu Gly
                    70
                                        75
Ala Gly Lys Ser Ser Trp Trp Val Ile Asn Pro Asp Ala Lys Pro Gly
```

85 90 Met Asn Pro Arg Arg Thr Arg Glu Arg Ser

100

```
<210> 38
<211> 60
<212> PRT
<213> Caenorhabditis elegans
<400> 38
Glu Ile Lys Leu Ser Asp Phe Lys His Gln Leu Phe Glu Leu Ile Ala
Pro Met Lys Trp Gly Thr Tyr Ser Val Lys Pro Gln Asp Tyr Val Phe
                               25
Arg Gln Leu Asn Asn Phe Gly Glu Ile Glu Val Ile Phe Asn Asp Asp
                           40
Gln Pro Leu Ser Lys Leu Glu Leu His Gly Thr Phe
<210> 39
<211> 2784
<212> DNA
<213> Caenorhabditis elegans
<400> 39
atqaaqctaa taqcaacttc tcttctagtt cccgacgagc acacaccgat gatgtcacca
                                                                      60
gtgaatacaa ctacaaagat tctacaacgg agtggtatta aaatggaaat cccgccatat
                                                                     120
                                                                     180
ttggatccag acagtcagga tgatgacccg gaagatggtg tcaactaccc ggatccagat
ttatttgaca caaaaaacac aaatatgacc gagtacgatt tggatgtgtt gaagcttgga
                                                                     240
aaaccagcag tagatgaagc acggaaaaag atcgaagttc ccgacgctag tgcgccgcca
                                                                     300
aacaaaattg tagaatattt gatgtattat agaacgttaa aagaaagtga actcatacaa
                                                                     360
ctqaatqcqt atcqqacaaa acqaaatcqa ttatcqttqa acttqqtcaa aaacaatatt
                                                                     420
qatcqaqaqt tcqaccaaaa agcttqcqaq tccctqqtqa aaaaattqaa gqataaqaaq
                                                                     480
aatgatctcc agaacctgat tgatgtggtt ctttcaaaaq gtacaaaata taccggttgc
                                                                     540
attacaattc caaggacact tgatggccgg ttacaggtcc acggaagaaa aggtttccct
                                                                     600
660
gtggaccact gcaagcacgc atttgaaatg aaaagtgaca tggtatgcgt gaatccctat
                                                                     720
cactacgaaa ttgtcattgg aactatgatt gttgggcaga gggatcatga caatcgagat
                                                                     780
atgccgccgc cacatcaacg ctaccacact ccaggtcggc aggatccagt tgacgatatg
                                                                     840
agtagattta taccaccage ttecattegt cegecteega tgaacatgca cacaaggeet
                                                                     900
cagoctatgo otcaacaatt goottoagtt ggogcaacgt ttgcccatco totcccacat
                                                                     960
caggogocac ataacccagg ggtttcacat cogtactcca ttgctccaca gacccattac
                                                                    1020
ccgttgaaca tgaacccaat tccgcaaatg ccgcaaatgc cacaaatgcc accacctctc
                                                                    1080
catcagggat atggaatgaa tgggccgagt tgctcttcag aaaacaacaa tccattccac
                                                                    1140
caaaatcacc attataatqa tattaqccat ccaaatcact attcctacqa ctqtqqtccq
                                                                    1200
aacttgtacg ggtttccaac tccttatccg gattttcacc atcctttcaa tcagcaacca
                                                                    1260
caccaqccqc cacaactatc acaaaaccat acqtcccaac aaqqcaqtca tcaaccaqqq
                                                                    1320
caccaaggtc aggtaccgaa tgatccacca atttcaagac cagtgttaca accatcaaca
                                                                    1380
gtcaccttgg acgtgttccg tcggtactgt agacagacat ttggaaatcg attttttgaa
                                                                    1440
ggagaaagtg aacaatccgg cgcaataatt cggtctagta acaaattcat tgaagaattt
                                                                    1500
gattcgccga tttgtggtgt gacagttgtt cgaccgcgga tgacagacgg tgaggttttg
                                                                    1560
gagaacatca tgccggaaga tgcaccatat catgacattt gcaagttcat tttgaggctc
                                                                    1620
acatcagaaa gtgtaacttt ctcaggagag gggccagaag ttagtgattt gaacgaaaaa
                                                                    1680
tggggaacaa ttgtgtacta tgagaaaaat ttgcaaattg gcgagaaaaa atgttcgaga
                                                                    1740
ggaaatttcc acgtggatgg cggattcatt tgctctgaga atcgttacag tctcggactt
                                                                    1800
gagccaaatc caattagaga accagtggcg tttaaagttc gtaaagcaat agtggatgga
                                                                    1860
attcgctttt cctacaaaaa agacgggagt gtttggcttc aaaaccgcat gaagtacccg
                                                                    1920
                                                                    1980
gtatttgtca cttctgggta tctcgacgag caatcaggag gcctaaagaa ggataaagtg
cacaaagttt acggatgtgc gtctatcaaa acgtttggct tcaacgtttc caaacaaatc
                                                                    2040
atcagagacg cgcttctttc caagcaaatg gcaacaatgt acttgcaagg aaaattgact
                                                                    2100
ccgatgaatt atatctacga gaagaagact caggaagagc tgcgaaggga agcaacacgc
                                                                    2160
```

accactgatt cattggccaa gtactgttgt gtccgtgtct cgttctgcaa aggatttgga

```
gaagcatacc cagaacgccc gtcaattcat gattgtccag tttggattga gttgaaaatc
                                                                      2280
                                                                      2340
aacattgcct acgatttcat ggattcaatc tgccagtaca taaccaactg cttcgagccg
                                                                      2400
ctaggaatgg aagattttgc aaaattggga atcaacgtca gtgatgacta aatgataact
tttttcactc accctactag atactgattt agtcttattc caaatcatcc aacgatatca
                                                                      2460
aactttttcc tttgaacttt gcatactatg ttatcacaag ttccaagcag tttcaataca
                                                                      2520
aacataggat atgttaacaa cttttgataa gaatcaagtt accaactgtt cattgtgagc
                                                                      2580
tttgagctgt atagaaggac aatgtatccc atacctcaat ctttaatagt catcagtcac
                                                                      2640
                                                                      2700
tggtcccgca ccaatttttt cgattcgcat atgtcatata ttgcaccgtg gcccttttta
                                                                      2760
ttgtaacttt taatatattt tcttcccaac ttgtgaatat gattgatgaa ccaccatttt
gagtaataaa tgtatttttt gtgg
                                                                      2784
<210> 40
<211> 796
<212> PRT
<213> Caenorhabditis elegans
<400> 40
Met Lys Leu Ile Ala Thr Ser Leu Leu Val Pro Asp Glu His Thr Pro
                                    10
Met Met Ser Pro Val Asn Thr Thr Lys Ile Leu Gln Arg Ser Gly
                                25
Ile Lys Met Glu Ile Pro Pro Tyr Leu Asp Pro Asp Ser Gln Asp Asp
                            40
Asp Pro Glu Asp Gly Val Asn Tyr Pro Asp Pro Asp Leu Phe Asp Thr
                        55
                                            60
Lys Asn Thr Asn Met Thr Glu Tyr Asp Leu Asp Val Leu Lys Leu Gly
                    70
                                        75
Lys Pro Ala Val Asp Glu Ala Arg Lys Lys Ile Glu Val Pro Asp Ala
               85
                                    90
Ser Ala Pro Pro Asn Lys Ile Val Glu Tyr Leu Met Tyr Tyr Arg Thr
           100
                                105
                                                    110
Leu Lys Glu Ser Glu Leu Ile Gln Leu Asn Ala Tyr Arg Thr Lys Arg
       115
                            120
                                                125
Asn Arg Leu Ser Leu Asn Leu Val Lys Asn Asn Ile Asp Arg Glu Phe
   130
                        135
                                            140
Asp Gln Lys Ala Cys Glu Ser Leu Val Lys Lys Leu Lys Asp Lys Lys
                    150
                                        155
                                                             160
Asn Asp Leu Gln Asn Leu Ile Asp Val Val Leu Ser Lys Gly Thr Lys
```

Tyr Thr Gly Cys Ile Thr Ile Pro Arg Thr Leu Asp Gly Arg Leu Gln Val His Gly Arg Lys Gly Phe Pro His Val Val Tyr Gly Lys Leu Trp Arg Phe Asn Glu Met Thr Lys Asn Glu Thr Arg His Val Asp His Cys Lys His Ala Phe Glu Met Lys Ser Asp Met Val Cys Val Asn Pro Tyr His Tyr Glu Ile Val Ile Gly Thr Met Ile Val Gly Gln Arg Asp His Asp Asn Arg Asp Met Pro Pro Pro His Gln Arg Tyr His Thr Pro Gly Arg Gln Asp Pro Val Asp Asp Met Ser Arg Phe Ile Pro Pro Ala Ser Ile Arg Pro Pro Pro Met Asn Met His Thr Arg Pro Gln Pro Met Pro Gln Gln Leu Pro Ser Val Gly Ala Thr Phe Ala His Pro Leu Pro His Gln Ala Pro His Asn Pro Gly Val Ser His Pro Tyr Ser Ile Ala Pro

```
Gln Thr His Tyr Pro Leu Asn Met Asn Pro Ile Pro Gln Met Pro Gln
            340
                               345
Met Pro Gln Met Pro Pro Pro Leu His Gln Gly Tyr Gly Met Asn Gly
                           360
       355
                                                365
Pro Ser Cys Ser Ser Glu Asn Asn Pro Phe His Gln Asn His His
                        375
                                            380
Tyr Asn Asp Ile Ser His Pro Asn His Tyr Ser Tyr Asp Cys Gly Pro
                   390
                                       395
Asn Leu Tyr Gly Phe Pro Thr Pro Tyr Pro Asp Phe His His Pro Phe
               405
                                   410
Asn Gln Gln Pro His Gln Pro Pro Gln Leu Ser Gln Asn His Thr Ser
                                425
Gln Gln Gly Ser His Gln Pro Gly His Gln Gly Gln Val Pro Asn Asp
                            440
Pro Pro Ile Ser Arg Pro Val Leu Gln Pro Ser Thr Val Thr Leu Asp
                       455
Val Phe Arg Arg Tyr Cys Arg Gln Thr Phe Gly Asn Arg Phe Phe Glu
                   470
                                       475
Gly Glu Ser Glu Gln Ser Gly Ala Ile Ile Arg Ser Ser Asn Lys Phe
               485
                                    490
Ile Glu Glu Phe Asp Ser Pro Ile Cys Gly Val Thr Val Val Arg Pro
                               505
Arg Met Thr Asp Gly Glu Val Leu Glu Asn Ile Met Pro Glu Asp Ala
                            520
Pro Tyr His Asp Ile Cys Lys Phe Ile Leu Arg Leu Thr Ser Glu Ser
                       535
Val Thr Phe Ser Gly Glu Gly Pro Glu Val Ser Asp Leu Asn Glu Lys
                   550
                                       555
Trp Gly Thr Ile Val Tyr Tyr Glu Lys Asn Leu Gln Ile Gly Glu Lys
               565
                                   570
Lys Cys Ser Arg Gly Asn Phe His Val Asp Gly Gly Phe Ile Cys Ser
                               585
Glu Asn Arg Tyr Ser Leu Gly Leu Glu Pro Asn Pro Ile Arg Glu Pro
                           600
Val Ala Phe Lys Val Arg Lys Ala Ile Val Asp Gly Ile Arg Phe Ser
                       615
                                            620
Tyr Lys Lys Asp Gly Ser Val Trp Leu Gln Asn Arg Met Lys Tyr Pro
                   630
                                        635
Val Phe Val Thr Ser Gly Tyr Leu Asp Glu Gln Ser Gly Gly Leu Lys
               645
                                    650
Lys Asp Lys Val His Lys Val Tyr Gly Cys Ala Ser Ile Lys Thr Phe
                               665
Gly Phe Asn Val Ser Lys Gln Ile Ile Arg Asp Ala Leu Leu Ser Lys
                           680
Gln Met Ala Thr Met Tyr Leu Gln Gly Lys Leu Thr Pro Met Asn Tyr
                       695
                                            700
Ile Tyr Glu Lys Lys Thr Gln Glu Glu Leu Arg Arg Glu Ala Thr Arg
                   710
Thr Thr Asp Ser Leu Ala Lys Tyr Cys Cys Val Arg Val Ser Phe Cys
                                   730
Lys Gly Phe Gly Glu Ala Tyr Pro Glu Arg Pro Ser Ile His Asp Cys
           740
                               745
Pro Val Trp Ile Glu Leu Lys Ile Asn Ile Ala Tyr Asp Phe Met Asp
                           760
Ser Ile Cys Gln Tyr Ile Thr Asn Cys Phe Glu Pro Leu Gly Met Glu
                       775
Asp Phe Ala Lys Leu Gly Ile Asn Val Ser Asp Asp
```

```
<211> 858
<212> PRT
<213> Caenorhabditis elegans
<400> 41
Met Gly Asp His His Asn Leu Thr Gly Leu Pro Gly Thr Ser Ile Pro
Pro Gln Phe Asn Tyr Ser Gln Pro Gly Thr Ser Thr Gly Gly Pro Leu
                                25
Tyr Gly Gly Lys Pro Ser His Gly Leu Glu Asp Ile Pro Asp Val Glu
                            40
Glu Tyr Glu Arg Asn Leu Leu Gly Ala Gly Ala Gly Phe Asn Leu Leu
                        55
Asn Val Gly Asn Met Ala Asn Val Pro Asp Glu His Thr Pro Met Met
                    70
                                        75
Ser Pro Val Asn Thr Thr Lys Ile Leu Gln Arg Ser Gly Ile Lys
                85
                                    90
Met Glu Ile Pro Pro Tyr Leu Asp Pro Asp Ser Gln Asp Asp Asp Pro
            100
                                105
                                                    110
Glu Asp Gly Val Asn Tyr Pro Asp Pro Asp Leu Phe Asp Thr Lys Asn
                            120
        115
                                                125
Thr Asn Met Thr Glu Tyr Asp Leu Asp Val Leu Lys Leu Gly Lys Pro
                        135
                                            140
Ala Val Asp Glu Ala Arg Lys Lys Ile Glu Val Pro Asp Ala Ser Ala
                    150
                                        155
Pro Pro Asn Lys Ile Val Glu Tyr Leu Met Tyr Tyr Arg Thr Leu Lys
                165
                                    170
                                                         175
Glu Ser Glu Leu Ile Gln Leu Asn Ala Tyr Arg Thr Lys Arg Asn Arg
            180
                                185
                                                    190
Leu Ser Leu Asn Leu Val Lys Asn Asn Ile Asp Arg Glu Phe Asp Gln
                            200
        195
                                                205
Lys Ala Cys Glu Ser Leu Val Lys Lys Leu Lys Asp Lys Lys Asn Asp
                        215
                                            220
Leu Gln Asn Leu Ile Asp Val Val Leu Ser Lys Gly Thr Lys Tyr Thr
                    230
                                        235
Gly Cys Ile Thr Ile Pro Arg Thr Leu Asp Gly Arg Leu Gln Val His
                                    250
                245
Gly Arg Lys Gly Phe Pro His Val Val Tyr Gly Lys Leu Trp Arg Phe
            260
                                                    270
                                265
Asn Glu Met Thr Lys Asn Glu Thr Arg His Val Asp His Cys Lys His
                            280
Ala Phe Glu Met Lys Ser Asp Met Val Cys Val Asn Pro Tyr His Tyr
                        295
                                            300
Glu Ile Val Ile Gly Thr Met Ile Val Gly Gln Arg Asp His Asp Asn
                    310
                                        315
Arg Asp Met Pro Pro Pro His Gln Arg Tyr His Thr Pro Gly Arg Gln
                                    330
                325
                                                         335
Asp Pro Val Asp Asp Met Ser Arg Phe Ile Pro Pro Ala Ser Ile Arg
                                345
            340
                                                    350
Pro Pro Met Asn Met His Thr Arg Pro Gln Pro Met Pro Gln Gln
                                              , 365
        355
                            360
Leu Pro Ser Val Gly Ala Thr Phe Ala His Pro Leu Pro His Gln Ala
                        375
                                            380
Pro His Asn Pro Gly Val Ser His Pro Tyr Ser Ile Ala Pro Gln Thr
                    390
                                        395
His Tyr Pro Leu Asn Met Asn Pro Ile Pro Gln Met Pro Gln Met Pro
                405
                                    410
```

<210> 41

Gln Met Pro Pro Pro Leu His Gln Gly Tyr Gly Met Asn Gly Pro Ser

```
425
Cys Ser Ser Glu Asn Asn Pro Phe His Gln Asn His His Tyr Asn
                            440
                                                 445
Asp Ile Ser His Pro Asn His Tyr Ser Tyr Asp Cys Gly Pro Asn Leu
                        455
                                            460
Tyr Gly Phe Pro Thr Pro Tyr Pro Asp Phe His His Pro Phe Asn Gln
                    470
                                        475
Gln Pro His Gln Pro Pro Gln Leu Ser Gln Asn His Thr Ser Gln Gln
                485
                                    490
Gly Ser His Gln Pro Gly His Gln Gly Gln Val Pro Asn Asp Pro Pro
            500
                                505
Ile Ser Arg Pro Val Leu Gln Pro Ser Thr Val Thr Leu Asp Val Phe
        515
                            520
Arg Arg Tyr Cys Arg Gln Thr Phe Gly Asn Arg Phe Phe Glu Gly Glu
                        535
                                             540
Ser Glu Gln Ser Gly Ala Ile Ile Arg Ser Ser Asn Lys Phe Ile Glu
                    550
                                        555
Glu Phe Asp Ser Pro Ile Cys Gly Val Thr Val Val Arg Pro Arg Met
                565
                                    570
Thr Asp Gly Glu Val Leu Glu Asn Ile Met Pro Glu Asp Ala Pro Tyr
                                585
His Asp Ile Cys Lys Phe Ile Leu Arg Leu Thr Ser Glu Ser Val Thr
        595
                            600
Phe Ser Gly Glu Gly Pro Glu Val Ser Asp Leu Asn Glu Lys Trp Gly
                        615
Thr Ile Val Tyr Tyr Glu Lys Asn Leu Gln Ile Gly Glu Lys Lys Cys
                    630
Ser Arg Gly Asn Phe His Val Asp Gly Gly Phe Ile Cys Ser Glu Asn
                645
                                    650
Arg Tyr Ser Leu Gly Leu Glu Pro Asn Pro Ile Arg Glu Pro Val Ala
                                665
Phe Lys Val Arg Lys Ala Ile Val Asp Gly Ile Arg Phe Ser Tyr Lys
                            680
Lys Asp Gly Ser Val Trp Leu Gln Asn Arg Met Lys Tyr Pro Val Phe
   690
                        695
                                            700
Val Thr Ser Gly Tyr Leu Asp Glu Gln Ser Gly Gly Leu Lys Lys Asp
                    710
                                        715
Lys Val His Lys Val Tyr Gly Cys Ala Ser Ile Lys Thr Phe Gly Phe
                725
                                    730
Asn Val Ser Lys Gln Ile Ile Arg Asp Ala Leu Leu Ser Lys Gln Met
            740
                                745
Ala Thr Met Tyr Leu Gln Gly Lys Leu Thr Pro Met Asn Tyr Ile Tyr
                            760
Glu Lys Lys Thr Gln Glu Glu Leu Arg Arg Glu Ala Thr Arg Thr Thr
   770
                        775
                                            780
Asp Ser Leu Ala Lys Tyr Cys Cys Val Arg Val Ser Phe Cys Lys Gly
                    790
                                        795
Phe Gly Glu Ala Tyr Pro Glu Arg Pro Ser Ile His Asp Cys Pro Val
                805
                                    810
Trp Ile Glu Leu Lys Ile Asn Ile Ala Tyr Asp Phe Met Asp Ser Ile
                                825
Cys Gln Tyr Ile Thr Asn Cys Phe Glu Pro Leu Gly Met Glu Asp Phe
                            840
Ala Lys Leu Gly Ile Asn Val Ser Asp Asp
    850
                        855
```

<210> 42 <211> 892

<400> 42 Met Gly Asp His His Asn Leu Thr Gly Leu Pro Gly Thr Ser Ile Pro Pro Gln Phe Asn Tyr Ser Gln Pro Gly Thr Ser Thr Gly Gly Pro Leu Tyr Gly Gly Lys Pro Ser His Gly Leu Glu Asp Ile Pro Asp Val Glu 40 Glu Tyr Glu Arg Asn Leu Leu Gly Ala Gly Ala Gly Phe Asn Leu Leu Asn Val Gly Asn Met Ala Asn Glu Phe Lys Pro Ile Ile Thr Leu Asp 75 Thr Lys Pro Pro Arg Asp Ala Asn Lys Ser Leu Ala Phe Asn Gly Gly 85 Leu Lys Leu Ile Thr Pro Lys Thr Glu Val Pro Asp Glu His Thr Pro 105 Met Met Ser Pro Val Asn Thr Thr Lys Ile Leu Gln Arg Ser Gly 115 120 125 Ile Lys Met Glu Ile Pro Pro Tyr Leu Asp Pro Asp Ser Gln Asp Asp 135 140 Asp Pro Glu Asp Gly Val Asn Tyr Pro Asp Pro Asp Leu Phe Asp Thr 150 155 Lys Asn Thr Asn Met Thr Glu Tyr Asp Leu Asp Val Leu Lys Leu Gly 165 170 Lys Pro Ala Val Asp Glu Ala Arg Lys Lys Ile Glu Val Pro Asp Ala 185 190 Ser Ala Pro Pro Asn Lys Ile Val Glu Tyr Leu Met Tyr Tyr Arg Thr 195 200 205 Leu Lys Glu Ser Glu Leu Ile Gln Leu Asn Ala Tyr Arg Thr Lys Arg 210 215 220 Asn Arg Leu Ser Leu Asn Leu Val Lys Asn Asn Ile Asp Arg Glu Phe 230 235 Asp Gln Lys Ala Cys Glu Ser Leu Val Lys Lys Leu Lys Asp Lys Lys 245 250 Asn Asp Leu Gln Asn Leu Ile Asp Val Val Leu Ser Lys Gly Thr Lys 265 Tyr Thr Gly Cys Ile Thr Ile Pro Arg Thr Leu Asp Gly Arg Leu Gln 275 280 Val His Gly Arg Lys Gly Phe Pro His Val Val Tyr Gly Lys Leu Trp 295 Arg Phe Asn Glu Met Thr Lys Asn Glu Thr Arg His Val Asp His Cys 310 315 Lys His Ala Phe Glu Met Lys Ser Asp Met Val Cys Val Asn Pro Tyr 325 330 His Tyr Glu Ile Val Ile Gly Thr Met Ile Val Gly Gln Arg Asp His 340 345 Asp Asn Arg Asp Met Pro Pro Pro His Gln Arg Tyr His Thr Pro Gly 355 360 Arg Gln Asp Pro Val Asp Asp Met Ser Arg Phe Ile Pro Pro Ala Ser 375 Ile Arg Pro Pro Pro Met Asn Met His Thr Arg Pro Gln Pro Met Pro 390 395 Gln Gln Leu Pro Ser Val Gly Ala Thr Phe Ala His Pro Leu Pro His 405 410 Gln Ala Pro His Asn Pro Gly Val Ser His Pro Tyr Ser Ile Ala Pro 425 Gln Thr His Tyr Pro Leu Asn Met Asn Pro Ile Pro Gln Met Pro Gln

```
435
                            440
Met Pro Gln Met Pro Pro Pro Leu His Gln Gly Tyr Gly Met Asn Gly
                        455
                                            460
Pro Ser Cys Ser Ser Glu Asn Asn Pro Phe His Gln Asn His His
                    470
                                        475
Tyr Asn Asp Ile Ser His Pro Asn His Tyr Ser Tyr Asp Cys Gly Pro
                485
                                    490
                                                        495
Asn Leu Tyr Gly Phe Pro Thr Pro Tyr Pro Asp Phe His His Pro Phe
            500
                                505
Asn Gln Gln Pro His Gln Pro Pro Gln Leu Ser Gln Asn His Thr Ser
                            520
Gln Gln Gly Ser His Gln Pro Gly His Gln Gly Gln Val Pro Asn Asp
                        535
                                            540
Pro Pro Ile Ser Arg Pro Val Leu Gln Pro Ser Thr Val Thr Leu Asp
                    550
                                        555
Val Phe Arg Arg Tyr Cys Arg Gln Thr Phe Gly Asn Arg Phe Phe Glu
                565
                                    570
Gly Glu Ser Glu Gln Ser Gly Ala Ile Ile Arg Ser Ser Asn Lys Phe
                                585
Ile Glu Glu Phe Asp Ser Pro Ile Cys Gly Val Thr Val Val Arg Pro
                            600
Arg Met Thr Asp Gly Glu Val Leu Glu Asn Ile Met Pro Glu Asp Ala
                        615
                                            620
Pro Tyr His Asp Ile Cys Lys Phe Ile Leu Arg Leu Thr Ser Glu Ser
                    630
                                        635
Val Thr Phe Ser Gly Glu Gly Pro Glu Val Ser Asp Leu Asn Glu Lys
                645
                                    650
Trp Gly Thr Ile Val Tyr Tyr Glu Lys Asn Leu Gln Ile Gly Glu Lys
                                665
Lys Cys Ser Arg Gly Asn Phe His Val Asp Gly Gly Phe Ile Cys Ser
                            680
Glu Asn Arg Tyr Ser Leu Gly Leu Glu Pro Asn Pro Ile Arg Glu Pro
                        695
Val Ala Phe Lys Val Arg Lys Ala Ile Val Asp Gly Ile Arg Phe Ser
                    710
                                        715
Tyr Lys Lys Asp Gly Ser Val Trp Leu Gln Asn Arg Met Lys Tyr Pro
                725
                                    730
Val Phe Val Thr Ser Gly Tyr Leu Asp Glu Gln Ser Gly Gly Leu Lys
            740
                                745
Lys Asp Lys Val His Lys Val Tyr Gly Cys Ala Ser Ile Lys Thr Phe
                            760
Gly Phe Asn Val Ser Lys Gln Ile Ile Arg Asp Ala Leu Leu Ser Lys
                        775
Gln Met Ala Thr Met Tyr Leu Gln Gly Lys Leu Thr Pro Met Asn Tyr
                    790
                                        795
Ile Tyr Glu Lys Lys Thr Gln Glu Glu Leu Arg Arg Glu Ala Thr Arg
                805
                                    810
Thr Thr Asp Ser Leu Ala Lys Tyr Cys Cys Val Arg Val Ser Phe Cys
                                825
Lys Gly Phe Gly Glu Ala Tyr Pro Glu Arg Pro Ser Ile His Asp Cys
                            840
Pro Val Trp Ile Glu Leu Lys Ile Asn Ile Ala Tyr Asp Phe Met Asp
                        855
                                            860
Ser Ile Cys Gln Tyr Ile Thr Asn Cys Phe Glu Pro Leu Gly Met Glu
                   870
                                        875
Asp Phe Ala Lys Leu Gly Ile Asn Val Ser Asp Asp
```

<210> 43 <211> 3499 <212> DNA <213> Caenorhabditis elegans

<400> 43 60 tgatctttca agccgaagca atcaagacct caaagccaat caactctact cacttttctt cagaacctta actttttgtg tcactttccc caaaaaccgt tcaagctgct gccttcactc 120 tcatcccctc ctcttactcc ttctttctcg tccgctacta ctgtatcttc tggacatcta 180 cctgtataca caccagtggc cagtcatctg ccattacaat ttcatcaatt gacacttctt 240 caacaacaac cgccgtcctc attcactccc gattcttcct catcctcaac atcgtcgtct 300 ttggctgaaa ttcccgaaga cgttatgatg gagatgctgg tagatcaggg aactgatgca 360 420 tegteateeg cetecaegte caceteatet gtttegagat teggagegga caegtteatg 480 aatacaccgg atgatgtgat gatgaatgat gatatggaac cgattcctcg tgatcggtgc aatacgtggc caatgcgtag gccgcaactc gaaccaccac tcaactcgag tcccattatt 540 600 catgaacaaa ttcctgaaga agatgctgac ctatacggga gcaatgagca atgtggacag 660 ctcggcggag catcttcaaa cgggtcgaca gcaatgcttc atactccaga tggaagcaat 720 tctcatcaga catcgtttct tcggagtttc agaatgtccg aatcgccaga cgataccgta tcgggaaaaa agacaacgac cagacggaac gcttggggaa atatgtcata tgctgaactt 780 atcactacag ccattatggc tagtccagag aaacggttaa ctcttgcaca agtttacgaa 840 900 tggatggtcc agaatgttcc atacttcagg gataagggag attcgaacag ttcagctgga 960 tggaagaact cgatccgtca caatctgtct cttcattctc gtttcatgcg aattcagaat 1020 gaaggagccg gaaagagctc gtggtgggtt attaatccag atgcaaagcc aggaatgaat ccacggcgta cacgtgaacg atccaatact attgagacga ctacaaaggc tcaactcgaa 1080 1140 aaatctcgcc gcggagccaa gaagaggata aaggagagag cattgatggg ctcccttcac tcgacactta atggaaattc gattgccgga tcgattcaaa cgatttctca cgatttgtat 1200 1260 gatgatgatc aatgcaagga gcatttgata aegttccatc atctttccgt ccccgaactc 1320 aatcgaacct ctcgattcct ggatcgtcgt ctcgtgtttc tccagctatt ggaagtgata 1380 tctatgatga tctagaattc ccatcatggg ttggcgaatc ggttccagca attccaagtg 1440 atattqttqa taqaactqat caaatqcqta tcgatqcaac tactcataqt tqgtgqagtt cagattaagc aggaqtcgaa gccgattaag acggaaccaa ttgctccacc accatcatac 1500 cacgagttga acagtgtccg tggatcgtgt gctcagaatc cacttcttcg aaatccaatt 1560 gtgccaagca ctaacttcaa gccaatgcca ctaccgggtg cctatggaaa ctatcaaaat 1620 ggtggaataa ctccaatcaa ttggctatca acatccaact catctccact gcctggaatt 1680 caatcgtgtg gaattgtagc tgcacagcat actgtcgctt cttcatcggc tcttccaatt 1740 gatttggaaa atctgacact tcccgatcag ccactgatgg atactatgga tgttgatgca 1800 ttgatcagac atgagctgag tcaagctgga gggcagcata ttcattttga tttgtaaatt 1860 ctcttcattt tgtttcccct ggtgttgttc gaaagagaga tagcaaagca gcgaggagtg 1920 aggtaagcag caataaaaat tttggatttt tttttggttt ttccagaaat aatcgatttt 1980 ctggaaaatt tcaaaaaaaa atcggaattt ttagttaatt atttgatgag aaaaaaaaat 2040 tagaaaacat aaggaaaaat gaaaagcgtt tttttttttc gaaaatttta gaattctcct 2100 acatttccaa taagggcctt agaactgcaa acaaacaaaa attggaattt tcgaatcaaa 2160 aagttcccga ataaaagtag ttcgaatatt aaaaagcatt taatttcctc tttaaaaaaaa 2220 ttgaataata gccgaaattt gcagattttt tttctgaaaa tcgaaaaacc aaaatttttt 2280 gattttttaa atttttttt tactttccag atagtaaaat cattagcact gaaaattatt 2340 tgaaaaaaa cttcaaatac aaattttgtt ttcgaaaaaa aaaatttaaa tatatttt 2400 cagaaatctt ccgtcttcat cttttcaaat ccctacctac acacactcaa cgatcatcac 2460 agccagacca toaatattot tocaaatttt gacgtcgtta attttttttc agttttttca 2520 aaaactctat tttctatttt ctgtcgtttg ttcccctttc tctcgtctaa ttccaacaca 2580 ttcatcccag tgacgtcgtg taataataat ataaaatacc tcttctctct ttcttcccct 2640 aatgcgaaat atcgaaaaac cgttgattat tacctctttt ttcttgtttt ttttttctct 2700 2760 ctctctctcc cgtcatccag gttcttcact ctttaaatgc tacctctatc ccatcttttt cgctgtaaat ttgtttcgca atcaaaactg ctaaaacaca ttccccaatc tgtctttttt 2820 aattgaattt ttcaaaaaat ttgatttctt gatttctctt gtaattcttt aattttcctc 2880 tttttttcc ccctggtagc aaatgtctag cgattctctt tctttttttg tttaactttc 2940 acatetggce gattegaate etcegtatae acacacacat agtaatetae etceaaaatt 3000 3060 ttactgaaag atgtgatccc ctctctgtct ccctctacaa aacattattt gtctgtttgt gtatattgcc accacgtcga ttttaaatta aaaccatcgt tttttcttct tttctacttt 3120 3180 tttctcgaaa aatttaacaa cacacaaaaa aatccttcaa aaaatctcag ttttaaatgg

```
tgtggcaata tatcggatcc ccctctacac cagaacagtc ttgcaatttc agagaatgat
                                                                     3240
titeagattt tteatateae aggeeeett ttittgettg ttitttete tacetetett
                                                                     3300
tottttcatt ctatttctct ctcttgtttt ctctctgtta tcctgtacat tttccttcca
                                                                     3360
attettetg getatttetg attttegagt teatattete taegteteae tttetetege
                                                                     3420
gccacgccc ctttttcgtc tccctccqcc cccaaatata tttgcgactg tatgatgatg
                                                                     3480
atgatgattt aataaaaat
                                                                     3499
<210> 44
<211> 2704
<212> DNA
<213> Caenorhabditis elegans
<400> 44
ttacacqtqq ccaatqcaac aatacatcta tcaqqaatcq tcaqcaacca ttccccatca
                                                                       60
ccatttaaat caacacaaca atccqtatca tccaatqcat cctcatcatc aattacctca
                                                                      120
tatgcaacaa cttcctcaac ctctattgaa tcttaacatg acqacgttaa catcttctqq
                                                                      180
cagttccgtg gccagttcca ttggaggcgg agctcaatgc tctccgtgcg cgtcgggctc
                                                                      240
ctcgaccgct gcaacaaatt cctctcaaca gcagcagacc gttggtcaaa tgcttgctgc
                                                                      300
ateggtgcct tgttcttcat ctggcatgac acttggaatg tcacttaatc tgtcacaagg
                                                                      360
cggtggtcca atgccggcaa aaaagaagcg ttgtcgtaag aagccaaccg atcaattggc
                                                                      420
acagaagaaa ccgaatccat ggggtgagga atcctattcg gatatcattg ccaaagcatt
                                                                      480
ggaatcggcg ccagacggaa ggcttaaact caatgagatt tatcaatggt tctctqataa
                                                                      540
tattccctac tttggagaac gatctagtcc cgaggaggcc gccggatgga agaactcgat
                                                                      600
ccgtcacaat ctgtctcttc attctcgttt catgcgaatt cagaatgaag gagccggaaa
                                                                      660
gagetegtgg tgggttatta atccagatge aaagecagga atgaatecae ggegtacaeg
                                                                      720
tgaacgatcc aatactattg agacgactac aaaggctcaa ctcgaaaaat ctcgccqcgg
                                                                      780
agccaagaag aggataaagg agagagcatt gatgggctcc cttcactcga cacttaatgg
                                                                      840
aaattcgatt gccggatcga ttcaaacgat ttctcacgat ttgtatgatg atgattcaat
                                                                      900
gcaaggagca tttgataacg ttccatcatc tttccgtccc cgaactcaat cgaacctctc
                                                                      960
gattcctgga tcgtcgtctc gtgtttctcc agctattgga agtgatatct atgatgatct
                                                                     1020
agaattccca tcatgggttg gcgaatcggt tccagcaatt ccaagtgata ttgttgatag
                                                                     1080
aactgatcaa atgcgtatcg atgcaactac tcatattggt ggagttcaga ttaagcagga
                                                                     1140
gtcgaagccg attaagacgg aaccaattgc tccaccacca tcataccacg agttgaacag
                                                                     1200
tgtccgtgga tcgtgtgctc agaatccact tcttcgaaat ccaattgtgc caagcactaa
                                                                     1260
cttcaagcca atgccactac cgggtgccta tggaaactat caaaatggtg gaataactcc
                                                                     1320
aatcaattgg ctatcaacat ccaactcatc tccactgcct ggaattcaat cgtgtggaat
                                                                     1380
tgtagctgca cagcatactg tcgcttcttc atcggctctt ccaattgatt tggaaaatct
                                                                     1440
gacacttccc gatcagccac tgatggatac tatggatgtt gatgcattga tcagacatga
                                                                     1500
gctgagtcaa gctggagggc agcatattca ttttgatttg taaattctct tcattttgtt
                                                                     1560
tcccctqqtq ttqttcqaaa qaqaqataqc aaaqcaqcqa qqaqtqaqaa atcttccqtc
                                                                     1620
ttcatctttt caaatcccta cctacacaca ctcaacqatc atcacaqcca qaccatcaat
                                                                     1680
attettecaa attttgacgt cgttaatttt ttttcagttt tttcaaaaac tetatttet
                                                                     1740
attttctgtc gtttgttccc ctttctctcg tctaattcca acacattcat cccagtgacg
                                                                     1800
tcgtgtaata ataatataaa atacctcttc tctctttctt cccctaatgc gaaatatcga
                                                                     1860
aaaaccgttg attattacct cttttttctt gtttttttt tctctctct tctcccgtca
                                                                     1920
tccaggttct tcactcttta aatgctacct ctatcccatc tttttcgctg taaatttgtt
                                                                     1980
tcgcaatcaa aactgctaaa acacattccc caatctqtct tttttaattq aatttttcaa
                                                                     2040
aaaatttgat ttcttgattt ctcttgtaat tctttaattt tcctcttttt tttccccctg
                                                                     2100
gtagcaaatg tctagcgatt ctctttcttt ttttgtttaa ctttcacatc tggccgattc
                                                                     2160
gaatcctccg tatacacaca cacatagtaa tctacctcca aaattttact gaaagatgtg
                                                                     2220
atcccctctc tgtctccctc tacaaaacat tatttgtctg tttgtgtata ttgccaccac
                                                                     2280
gtcgatttta aattaaaacc atcgtttttt cttcttttct actttttct cgaaaaattt
                                                                     2340
aacaacaca aaaaaaatcc ttcaaaaaat ctcagtttta aatggtgtgg caatatatcg
                                                                     2400
gatccccctc tacaccagaa cagtcttgca atttcagaga atgattttca gatttttcat
                                                                     2460
atcacaggee coettittt gettgtttt ttetetacet etettett teattetatt
                                                                     2520
tetetetett gttttetete tgttateetg taeattttee tteeaattet ttetggetat
                                                                     2580
ttctgatttt cgagttcata ttctctacgt ctcactttct ctcgcgccac gcccctttt
                                                                     2640
tcgtctccct ccgcccccaa atatatttgc gactgtatga tgatgatgat gatttaataa
                                                                     2700
aaat
                                                                     2704
```

```
<211> 510
<212> PRT
<213> Caenorhabditis elegans
<400> 45
Met Met Glu Met Leu Val Asp Gln Gly Thr Asp Ala Ser Ser Ser Ala
1
                                     10
Ser Thr Ser Thr Ser Ser Val Ser Arg Phe Gly Ala Asp Thr Phe Met
            20
                                 25
Asn Thr Pro Asp Asp Val Met Met Asn Asp Asp Met Glu Pro Ile Pro
                            40
Arg Asp Arg Cys Asn Thr Trp Pro Met Arg Arg Pro Gln Leu Glu Pro
                        55
Pro Leu Asn Ser Ser Pro Ile Ile His Glu Gln Ile Pro Glu Glu Asp
                    70
Ala Asp Leu Tyr Gly Ser Asn Glu Gln Cys Gly Gln Leu Gly Gly Ala
                85
                                     90
Ser Ser Asn Gly Ser Thr Ala Met Leu His Thr Pro Asp Gly Ser Asn
            100
                                 105
                                                     110
Ser His Gln Thr Ser Phe Pro Ser Asp Phe Arg Met Ser Glu Ser Pro
        115
                            120
Asp Asp Thr Val Ser Gly Lys Lys Thr Thr Thr Arg Arg Asn Ala Trp
                        135
                                             140
Gly Asn Met Ser Tyr Ala Glu Leu Ile Thr Thr Ala Ile Met Ala Ser
                    150
                                         155
Pro Glu Lys Arg Leu Thr Leu Ala Gln Val Tyr Glu Trp Met Val Gln
                165
                                     170
Asn Val Pro Tyr Phe Arg Asp Lys Gly Asp Ser Asn Ser Ser Ala Gly
            180
                                185
Trp Lys Asn Ser Ile Arg His Asn Leu Ser Leu His Ser Arg Phe Met
        195
                            200
Arg Ile Gln Asn Glu Gly Ala Gly Lys Ser Ser Trp Trp Val Ile Asn
                        215
                                             220
Pro Asp Ala Lys Pro Gly Met Asn Pro Arg Arg Thr Arg Glu Arg Ser
                    230
                                         235
Asn Thr Ile Glu Thr Thr Lys Ala Gln Leu Glu Lys Ser Arg Arg
                245
                                     250
Gly Ala Lys Lys Arg Ile Lys Glu Arg Ala Leu Met Gly Ser Leu His
            260
                                265
                                                     270
Ser Thr Leu Asn Gly Asn Ser Ile Ala Gly Ser Ile Gln Thr Ile Ser
        275
                            280
His Asp Leu Tyr Asp Asp Asp Ser Met Gln Gly Ala Phe Asp Asn Val
                        295
                                             300
Pro Ser Ser Phe Arg Pro Arg Thr Gln Ser Asn Leu Ser Ile Pro Gly
                    310
                                         315
                                                             320
Ser Ser Ser Arg Val Ser Pro Ala Ile Gly Ser Asp Ile Tyr Asp Asp
                325
                                     330
Leu Glu Phe Pro Ser Trp Val Gly Glu Ser Val Pro Ala Ile Pro Ser
            340
                                 345
Asp Ile Val Asp Arg Thr Asp Gln Met Arg Ile Asp Ala Thr Thr His
                            360
Ile Gly Gly Val Gln Ile Lys Gln Glu Ser Lys Pro Ile Lys Thr Glu
    370
                        375
Pro Ile Ala Pro Pro Pro Ser Tyr His Glu Leu Asn Ser Val Arg Gly
                    390
                                         395
                                                             400
Ser Cys Ala Gln Asn Pro Leu Leu Arg Asn Pro Ile Val Pro Ser Thr
Asn Phe Lys Pro Met Pro Leu Pro Gly Ala Tyr Gly Asn Tyr Gln Asn
```

<210> 45

```
420
                                425
Gly Gly Ile Thr Pro Ile Asn Trp Leu Ser Thr Ser Asn Ser Ser Pro
                            440
                                                445
Leu Pro Gly Ile Gln Ser Cys Gly Ile Val Ala Ala Gln His Thr Val
                        455
                                            460
Ala Ser Ser Ser Ala Leu Pro Ile Asp Leu Glu Asn Leu Thr Leu Pro
                    470
                                        475
Asp Gln Pro Leu Met Asp Thr Met Asp Val Asp Ala Leu Ile Arg His
                485
                                    490
Glu Leu Ser Gln Ala Gly Gly Gln His Ile His Phe Asp Leu
            500
                                505
<210> 46
<211> 509
<212> PRT
<213> Caenorhabditis elegans
<400> 46
Met Gln Gln Tyr Ile Tyr Gln Glu Ser Ser Ala Thr Ile Pro His His
His Leu Asn Gln His Asn Asn Pro Tyr His Pro Met His Pro His His
            20
Gln Leu Pro His Met Gln Gln Leu Pro Gln Pro Leu Leu Asn Leu Asn
Met Thr Thr Leu Thr Ser Ser Gly Ser Ser Val Ala Ser Ser Ile Gly
                        55
Gly Gly Ala Gln Cys Ser Pro Cys Ala Ser Gly Ser Ser Thr Ala Ala
                    70
Thr Asn Ser Ser Gln Gln Gln Thr Val Gly Gln Met Leu Ala Ala
Ser Val Pro Cys Ser Ser Ser Gly Met Thr Leu Gly Met Ser Leu Asn
            100
                                105
```

Leu Ser Gln Gly Gly Gly Pro Met Pro Ala Lys Lys Lys Arg Cys Arg 120 Lys Lys Pro Thr Asp Gln Leu Ala Gln Lys Lys Pro Asn Pro Trp Gly 130 135 Glu Glu Ser Tyr Ser Asp Ile Ile Ala Lys Ala Leu Glu Ser Ala Pro 150 Asp Gly Arg Leu Lys Leu Asn Glu Ile Tyr Gln Trp Phe Ser Asp Asn 165 170 Ile Pro Tyr Phe Gly Glu Arg Ser Ser Pro Glu Glu Ala Ala Gly Trp 185 Lys Asn Ser Ile Arg His Asn Leu Ser Leu His Ser Arg Phe Met Arg 200 Ile Gln Asn Glu Gly Ala Gly Lys Ser Ser Trp Trp Val Ile Asn Pro 215 220 Asp Ala Lys Pro Gly Met Asn Pro Arg Arg Thr Arg Glu Arg Ser Asn 230 235 Thr Ile Glu Thr Thr Lys Ala Gln Leu Glu Lys Ser Arg Arg Gly 245 250 Ala Lys Lys Arg Ile Lys Glu Arg Ala Leu Met Gly Ser Leu His Ser 265 Thr Leu Asn Gly Asn Ser Ile Ala Gly Ser Ile Gln Thr Ile Ser His 280 Asp Leu Tyr Asp Asp Asp Ser Met Gln Gly Ala Phe Asp Asn Val Pro 295 300 Ser Ser Phe Arg Pro Arg Thr Gln Ser Asn Leu Ser Ile Pro Gly Ser 310

```
Ser Ser Arg Val Ser Pro Ala Ile Gly Ser Asp Ile Tyr Asp Asp Leu
                325
                                     330
                                                          335
Glu Phe Pro Ser Trp Val Gly Glu Ser Val Pro Ala Ile Pro Ser Asp
                                 345
Ile Val Asp Arg Thr Asp Gln Met Arg Ile Asp Ala Thr Thr His Ile
                             360
Gly Gly Val Gln Ile Lys Gln Glu Ser Lys Pro Ile Lys Thr Glu Pro
                        375
                                             380
Ile Ala Pro Pro Pro Ser Tyr His Glu Leu Asn Ser Val Arg Gly Ser
                    390
                                         395
Cys Ala Gln Asn Pro Leu Leu Arg Asn Pro Ile Val Pro Ser Thr Asn
                405
                                     410
Phe Lys Pro Met Pro Leu Pro Gly Ala Tyr Gly Asn Tyr Gln Asn Gly
                                 425
Gly Ile Thr Pro Ile Asn Trp Leu Ser Thr Ser Asn Ser Ser Pro Leu
        435
                             440
                                                 445
Pro Gly Ile Gln Ser Cys Gly Ile Val Ala Ala Gln His Thr Val Ala
                        455
Ser Ser Ser Ala Leu Pro Ile Asp Leu Glu Asn Leu Thr Leu Pro Asp
                    470
                                         475
Gln Pro Leu Met Asp Thr Met Asp Val Asp Ala Leu Ile Arg His Glu
                485
                                     490
Leu Ser Gln Ala Gly Gly Gln His Ile His Phe Asp Leu
                                 505
```

<210> 47 <211> 3504 <212> DNA

<213> Caenorhabditis elegans

<400> 47

eggaagecat ggagetegag atetgattge tggacaegga eggaaeteeg aegtateteg 60 cagatgcatg ttaacatttt acatccacaa ctgcaaacga tggtcgagca gtggcaaatg 120 cgagaacgcc catcgctgga gaccgagaat ggcaaaggat cgctgctcct ggaaaatgaa 180 ggtgtcgcag atatcatcac tatgtgtcca ttcggagaag ttattagtgt agtatttccg 240 tggtttcttg caaatgtgcg aacatcgcta gaaatcaagc tatcagattt caaacatcaa 300 cttttcgaat tgattgctcc gatgaagtgg ggaacatatt ccgtaaagcc acaggattat 360 gtgttcagac agttgaataa tttcggcgaa attgaagtta tatttaacga cgatcaaccc 420 ctgtcgaaat tagagctcca cggcactttc ccaatgcttt ttctctacca acctgatgga 480 ataaacaggg ataaagaatt aatgagtgat ataagtcatt gtctaggata ctcactggat 540 aaactggaag agagcetega tgaggaacte egteaattte gtgettetet etgggetegt 600 acgaagaaaa cgtgcttgac acgtggactt gagggtacca gtcactacgc gttccccqaa 660 gaacagtact tgtgtgttgg tgaatcgtgc ccgaaagatt tggaatcaaa agtcaaqqct 720 gccaagctga gttatcagat gttttggaga aaacgtaaag cggaaatcaa tggagtttgc 780 gagaaaatga tgaagattca aattgaattc aatccgaacg aaactccgaa atctctgctt 840 cacacgtttc tctacgaaat gcgaaaattg gatgtatacg ataccgatga tcctgcagat 900 gaaggatggt ttcttcaatt ggctggacgt accacgtttg ttacaaatcc agatgtcaaa 960 cttacgtctt atgatggtgt ccgttcggaa ctggaaagct atcgatgccc tggattcgtt 1020 gttcgccgac aatcactagt cctcaaagac tattgtcgcc caaaaccact ctacgaacca 1080 cattatgtga gagcacacga acgaaaactt gctctaqacg tgctcagcgt gtctatagat 1140 agcacaccaa aacagagcaa gaacagtqac atggttatga ctgattttcg tccgacagct 1200 tcactcaaac aagtttcact ttgggacctt gacgcgaatc ttatgatacg gcctgtgaat 1260 atttctggat tcgatttccc ggccgacgtg gatatgtacg ttcgaatcga attcagtgta 1320 tatgtgggga cactgacgct ggcatcaaaa tctacaacaa aagtgaatgc tcaatttgca 1380 aaatggaata aggaaatgta cacttttgat ctatacatga aggatatgcc accatctgca 1440 gtactcagca ttcgtgtttt gtacggaaaa gtgaaattaa aaagtgaaga attcgaagtt 1500 ggttgggtaa atatgtccct aaccgattgg agagatgaac tacgacaagg acaattttta 1560 ttccatctgt gggctcctga accgactgcc aatcgtagta ggatcggaga aaatggagca 1620

```
aggataggca ccaacgcagc ggttacaatt gaaatctcaa gttatggtgg tagagttcga
                                                                      1680
atgeegagte aaggaeaata cacatatete gteaageace gaagtaettg gaeggaaact
                                                                      1740
ttgaatatta tgggtgatga ctatgagtcg tgtatcagag atccaggata taagaagctt
                                                                      1800
cagatgcttg tcaagaagca tgaatctgga attgtattag aggaagatga acaacgtcat
                                                                      1860
gtctggatgt ggaggagata cattcaaaag caggagcctg atttgctcat tgtgctctcc
                                                                      1920
gaactcgcat ttgtgtggac tgatcgtgag aacttttccg agctctatgt gatgcttgaa
                                                                      1980
aaatggaaac cgccgagtgt ggcagccgcg ttgactttgc ttggaaaacg ttgcacggat
                                                                      2040
cgtgtgattc gaaagtttgc agtggagaag ttgaatgagc agctgagccc ggtcacattc
                                                                      2100
catcttttca tattgcctct catacaggcg ttgaagtacg aaccgcgtgc tcaatcggaa
                                                                      2160
gttggaatga tgctcttgac tagagctctc tgcgattatc gaattggaca tcgacttttc
                                                                      2220
tggctgctcc gtgcagagat tgctcgtttg agagattgtg atctgaaaag tgaagaatat
                                                                      2280
egeegtatet caettetgat ggaagettae eteegtggaa atgaagagea cateaagate
                                                                      2340
atcacccgac aagttgacat ggttgatgag ctcacacgaa tcagcactct tgtcaaagga
                                                                      2400
atgccaaaag atgttgctac gatgaaactg cgtgacgagc ttcgatcgat tagtcataaa
                                                                      2460
atggaaaata tggattctcc actggatcct gtgtacaaac tgggtgaaat gataatcgac
                                                                      2520
aaagccatcg tcctaggaag tgcaaaacgt ccgttaatgc ttcactggaa gaacaaaaat
                                                                      2580
ccaaagagtg acctgcacct tccgttctgt gcaatgatct tcaagaatgg agacgatctt
                                                                      2640
cgccaggaca tgcttgttct tcaagttctc gaagttatgg ataacatctg gaaggctgca
                                                                      2700
aacattgatt gctgtttgaa cccgtacgca gttcttccaa tgggagaaat gattggaatt
                                                                      2760
attgaagttg tgcctaattg taaaacaata ttcgagattc aagttggaac aggattcatg
                                                                      2820
aatacagcag ttcggagtat tgatccttcg tttatgaata agtggattcg gaaacaatgc
                                                                      2880
ggaattgaag atgaaaagaa gaaaagcaaa aaggactcta cgaaaaatcc catcgaaaag
                                                                      2940
aagattgata atactcaagc catgaagaaa tattttgaaa gtgtcgatcg attcctatac
                                                                      3000
tcgtgtgttg gatattcagt tgccacgtac ataatgggaa tcaaggatcg tcacagtgat
                                                                      3060
aatctgatgc tcactgaaga tggaaaatat gtccacattg atttcggtca cattttggga
                                                                      3120
cacggaaaga ccaaacttgg qatccaqcga qatcqtcaac cqtttattct aaccgaacac
                                                                      3180
tttatgacag tgattcgatc qqqtaaatct qtqqatqqaa attcqcatqa qctacaaaaa
                                                                      3240
ttcaaaacgt tatgcgtcga agcctacgaa gtaatgtgga ataatcgaga tttgttcgtt
                                                                      3300
tecttgttca cettgatget eggaatggag ttgeetgage tgtegaegaa ageggatttg
                                                                      3360
gatcatttga agaaaaccct cttctgcaat ggagaaagca aagaagaagc gagaaagttt
                                                                      3420
ttcgctggaa tctacgaaga agccttcaat ggatcatggt ctaccaaaac gaattggctc
                                                                      3480
ttccacgcag tcaaacacta ctga
                                                                      3504
```

```
<210> 48
<211> 1167
<212> PRT
```

<213> Caenorhabditis elegans

<400> 48

Arg Lys Pro Trp Ser Ser Arg Ser Asp Cys Trp Thr Arg Thr Glu Leu 1 10 Arg Arg Ile Ser Gln Met His Val Asn Ile Leu His Pro Gln Leu Gln 20 25 30 Thr Met Val Glu Gln Trp Gln Met Arg Glu Arg Pro Ser Leu Glu Thr 40 Glu Asn Gly Lys Gly Ser Leu Leu Glu Asn Glu Gly Val Ala Asp 55 Ile Ile Thr Met Cys Pro Phe Gly Glu Val Ile Ser Val Val Phe Pro 70 75 Trp Phe Leu Ala Asn Val Arg Thr Ser Leu Glu Ile Lys Leu Ser Asp 90 Phe Lys His Gln Leu Phe Glu Leu Ile Ala Pro Met Lys Trp Gly Thr 105 110 Tyr Ser Val Lys Pro Gln Asp Tyr Val Phe Arg Gln Leu Asn Asn Phe 115 120 Gly Glu Ile Glu Val Ile Phe Asn Asp Asp Gln Pro Leu Ser Lys Leu 140 Glu Leu His Gly Thr Phe Pro Met Leu Phe Leu Tyr Gln Pro Asp Gly 145 150 160 155

```
Ile Asn Arg Asp Lys Glu Leu Met Ser Asp Ile Ser His Cys Leu Gly
                165
                                    170
Tyr Ser Leu Asp Lys Leu Glu Glu Ser Leu Asp Glu Glu Leu Arg Gln
            180
                                185
                                                     190
Phe Arg Ala Ser Leu Trp Ala Arg Thr Lys Lys Thr Cys Leu Thr Arg
        195
                            200
                                                 205
Gly Leu Glu Gly Thr Ser His Tyr Ala Phe Pro Glu Glu Gln Tyr Leu
                        215
                                             220
Cys Val Gly Glu Ser Cys Pro Lys Asp Leu Glu Ser Lys Val Lys Ala
225
                    230
                                        235
Ala Lys Leu Ser Tyr Gln Met Phe Trp Arg Lys Arg Lys Ala Glu Ile
                245
                                    250
Asn Gly Val Cys Glu Lys Met Met Lys Ile Gln Ile Glu Phe Asn Pro
            260
                                265
                                                     270
Asn Glu Thr Pro Lys Ser Leu Leu His Thr Phe Leu Tyr Glu Met Arg
                            280
                                                 285
Lys Leu Asp Val Tyr Asp Thr Asp Asp Pro Ala Asp Glu Gly Trp Phe
                        295
                                             300
Leu Gln Leu Ala Gly Arg Thr Thr Phe Val Thr Asn Pro Asp Val Lys
305
                    310
                                        315
Leu Thr Ser Tyr Asp Gly Val Arg Ser Glu Leu Glu Ser Tyr Arg Cys
                325
                                    330
Pro Gly Phe Val Val Arg Arg Gln Ser Leu Val Leu Lys Asp Tyr Cys
            340
                                345
                                                     350
Arg Pro Lys Pro Leu Tyr Glu Pro His Tyr Val Arg Ala His Glu Arg
        355
                            360
Lys Leu Ala Leu Asp Val Leu Ser Val Ser Ile Asp Ser Thr Pro Lys
                        375
                                             380
Gln Ser Lys Asn Ser Asp Met Val Met Thr Asp Phe Arg Pro Thr Ala
                    390
                                        395
Ser Leu Lys Gln Val Ser Leu Trp Asp Leu Asp Ala Asn Leu Met Ile
                405
                                    410
Arg Pro Val Asn Ile Ser Gly Phe Asp Phe Pro Ala Asp Val Asp Met
            420
                                425
                                                     430
Tyr Val Arg Ile Glu Phe Ser Val Tyr Val Gly Thr Leu Thr Leu Ala
        435
                            440
Ser Lys Ser Thr Thr Lys Val Asn Ala Gln Phe Ala Lys Trp Asn Lys
    450
                        455
                                            460
Glu Met Tyr Thr Phe Asp Leu Tyr Met Lys Asp Met Pro Pro Ser Ala
                    470
                                        475
Val Leu Ser Ile Arg Val Leu Tyr Gly Lys Val Lys Leu Lys Ser Glu
                485
                                    490
Glu Phe Glu Val Gly Trp Val Asn Met Ser Leu Thr Asp Trp Arg Asp
            500
                                505
                                                     510
Glu Leu Arg Gln Gly Gln Phe Leu Phe His Leu Trp Ala Pro Glu Pro
        515
                            520
                                                525
Thr Ala Asn Arg Ser Arg Ile Gly Glu Asn Gly Ala Arg Ile Gly Thr
                        535
                                            540
Asn Ala Ala Val Thr Ile Glu Ile Ser Ser Tyr Gly Gly Arg Val Arg
                    550
                                        555
Met Pro Ser Gln Gly Gln Tyr Thr Tyr Leu Val Lys His Arg Ser Thr
                565
                                    570
Trp Thr Glu Thr Leu Asn Ile Met Gly Asp Asp Tyr Glu Ser Cys Ile
                                585
Arg Asp Pro Gly Tyr Lys Lys Leu Gln Met Leu Val Lys Lys His Glu
        595
                            600
                                                605
Ser Gly Ile Val Leu Glu Glu Asp Glu Gln Arg His Val Trp Met Trp
                                            620
                        615
Arg Arg Tyr Ile Gln Lys Gln Glu Pro Asp Leu Leu Ile Val Leu Ser
```

```
630
                                     635
625
Glu Leu Ala Phe Val Trp Thr Asp Arg Glu Asn Phe Ser Glu Leu Tyr
           645
                                650
Val Met Leu Glu Lys Trp Lys Pro Pro Ser Val Ala Ala Ala Leu Thr
          660
                             665
Leu Leu Gly Lys Arg Cys Thr Asp Arg Val Ile Arg Lys Phe Ala Val
                        680
                                           685
Glu Lys Leu Asn Glu Gln Leu Ser Pro Val Thr Phe His Leu Phe Ile
                     695
                                        700
Leu Pro Leu Ile Gln Ala Leu Lys Tyr Glu Pro Arg Ala Gln Ser Glu
                 710
Val Gly Met Met Leu Leu Thr Arg Ala Leu Cys Asp Tyr Arg Ile Gly
              725
                                 730
His Arg Leu Phe Trp Leu Leu Arg Ala Glu Ile Ala Arg Leu Arg Asp
                             745
Cys Asp Leu Lys Ser Glu Glu Tyr Arg Arg Ile Ser Leu Leu Met Glu
                         760
Ala Tyr Leu Arg Gly Asn Glu Glu His Ile Lys Ile Ile Thr Arg Gln
                     775
Val Asp Met Val Asp Glu Leu Thr Arg Ile Ser Thr Leu Val Lys Gly
                 790
                                    795
Met Pro Lys Asp Val Ala Thr Met Lys Leu Arg Asp Glu Leu Arg Ser
             805
                                 810
Ile Ser His Lys Met Glu Asn Met Asp Ser Pro Leu Asp Pro Val Tyr
                             825
Lys Leu Gly Glu Met Ile Ile Asp Lys Ala Ile Val Leu Gly Ser Ala
                         840
Lys Arg Pro Leu Met Leu His Trp Lys Asn Lys Asn Pro Lys Ser Asp
                     855
                                        860
Leu His Leu Pro Phe Cys Ala Met Ile Phe Lys Asn Gly Asp Asp Leu
                 870
                                    875
Arg Gln Asp Met Leu Val Leu Gln Val Leu Glu Val Met Asp Asn Ile
                                 890
Trp Lys Ala Ala Asn Ile Asp Cys Cys Leu Asn Pro Tyr Ala Val Leu
                             905
Pro Met Gly Glu Met Ile Gly Ile Ile Glu Val Val Pro Asn Cys Lys
                         920
Thr Ile Phe Glu Ile Gln Val Gly Thr Gly Phe Met Asn Thr Ala Val
                      935
                                         940
Arg Ser Ile Asp Pro Ser Phe Met Asn Lys Trp Ile Arg Lys Gln Cys
                                    955
Gly Ile Glu Asp Glu Lys Lys Lys Ser Lys Lys Asp Ser Thr Lys Asn
                                 970
Pro Ile Glu Lys Lys Ile Asp Asn Thr Gln Ala Met Lys Lys Tyr Phe
                             985
Glu Ser Val Asp Arg Phe Leu Tyr Ser Cys Val Gly Tyr Ser Val Ala
                         1000
                                            1005
Thr Tyr Ile Met Gly Ile Lys Asp Arg His Ser Asp Asn Leu Met Leu
                     1015
                                        1020
Thr Glu Asp Gly Lys Tyr Val His Ile Asp Phe Gly His Ile Leu Gly
                 1030
                                    1035
His Gly Lys Thr Lys Leu Gly Ile Gln Arg Asp Arg Gln Pro Phe Ile
              1045
                                1050
Leu Thr Glu His Phe Met Thr Val Ile Arg Ser Gly Lys Ser Val Asp
          1060
                             1065 1070
Gly Asn Ser His Glu Leu Gln Lys Phe Lys Thr Leu Cys Val Glu Ala
  1075 1080 1085
Tyr Glu Val Met Trp Asn Asn Arg Asp Leu Phe Val Ser Leu Phe Thr
   1090
                      1095
                                         1100
```

```
1115
1105
                    1110
                                                              112
Asp His Leu Lys Lys Thr Leu Phe Cys Asn Gly Glu Ser Lys Glu Glu
                                     1130
                1125
Ala Arg Lys Phe Phe Ala Gly Ile Tyr Glu Glu Ala Phe Asn Gly Ser
            1140
                                 1145
Trp Ser Thr Lys Thr Asn Trp Leu Phe His Ala Val Lys His Tyr
                                                 1165
        1155
                             1160
<210> 49
<211> 23
<212> DNA
<213> Artificial Sequence
<220>
<223> Probe/primer derived from C. elegans
<400> 49
                                                                         23
ggaaatattt taggccagat gcg
<210> 50
<211> 20
<212> DNA
<213> Artificial Sequence
<220>
<223> Probe/primer derived from C. elegans
<400> 50
                                                                         20
cggacagtcc tgaatacacc
<210> 51
<211> 28
<212> DNA
<213> Artificial Sequence
<220>
<223> Probe/primer derived from C. elegans
<400> 51
                                                                         28
tctcgttgtt tgccgtcgga tgtctgcc
<210> 52
<211> 3017
<212> DNA
<213> Caenorhabditis elegans
<400> 52
gtaatcaaat tgtaaaggaa aaatattaat agtcagagta cacataaatg ggtgatcatc
                                                                         60
ataatttaac gggccttccc ggtacctcca tcccgccaca gttcaactat tctcagcccg
                                                                        120
gtaccagcac cggaggcccg ctttatggtg gaaaaccttc tcatggattg gaagatattc
                                                                        180
ctgatgtaga ggaatatgag aggaacctgc tcggggctgg agcaggtttt aatctgctca
                                                                        240
atgtaggaaa tatggctaat gttcccgacg agcacacacc gatgatgtca ccagtgaata
                                                                        300
caactacaaa gattctacaa cggagtggta ttaaaatgga aatcccgcca tatttggatc
                                                                        360
cagacagtca ggatgatgac ccggaagatg gtgtcaacta cccggatcca gatttatttg
                                                                        420
acacaaaaaa cacaaatatg accgagtacg atttggatgt gttgaagctt ggaaaaccag
                                                                        480
cagtagatga agcacggaaa aagatcgaag ttcccgacgc tagtgcgccg ccaaacaaaa
                                                                        540
ttgtagaata tttgatgtat tatagaacgt taaaagaaag tgaactcata caactgaatg
                                                                        600
```

Leu Met Leu Gly Met Glu Leu Pro Glu Leu Ser Thr Lys Ala Asp Leu

```
cgtatcggac aaaacgaaat cgattatcgt tgaacttggt caaaaacaat attgatcgag
                                                                       660
agttcgacca aaaagcttgc gagtccctgg tgaaaaaatt gaaqgataag aagaatgatc
                                                                       720
tccagaacct gattgatgtg gttctttcaa aaggtacaaa atataccggt tgcattacaa
                                                                       780
ttccaaggac acttgatggc cqqttacaqq tccacqqaaq aaaaqqtttc cctcacqtaq
                                                                       840
tctatggcaa actgtggagg tttaatgaaa tgacaaaaaa cgaaacgcgt catgtggacc
                                                                       900
actgcaagca cgcatttgaa atgaaaagtg acatggtatg cgtgaatccc tatcactacg
                                                                       960
aaattqtcat tqqaactatq attqttqqqc aqaqqqatca tqacaatcqa qatatqccqc
                                                                      1020
cgccacatca acgctaccac actccaggtc ggcaggatcc agttgacgat atgagtagat
                                                                      1080
ttataccacc agettecatt cgtccgcctc cgatqaacat gcacacaagg cctcaqccta
                                                                      1140
tgcctcaaca attgccttca gttggcgcaa cgtttgccca tcctctccca catcaggcgc
                                                                      1200
cacataaccc aggggtttca catccgtact ccattgctcc acagacccat tacccgttga
                                                                      1260
acatgaaccc aattccgcaa atgccgcaaa tgccacaaat gccaccacct ctccatcagg
                                                                      1320
gatatggaat gaatgggccg agttgctctt cagaaaacaa caatccattc caccaaaatc
                                                                      1380
accattataa tgatattagc catccaaatc actattccta cgactgtggt ccgaacttgt
                                                                      1440
acgggtttcc aactccttat ccggattttc accatccttt caatcagcaa ccacaccagc
                                                                      1500
cgccacaact atcacaaaac catacgtccc aacaaggcag tcatcaacca gggcaccaag
                                                                      1560
gtcaggtacc gaatgatcca ccaatttcaa gaccagtgtt acaaccatca acagtcacct
                                                                      1620
tggacgtgtt ccgtcggtac tgtagacaga catttggaaa tcgatttttt gaaggagaaa
                                                                      1680
gtgaacaatc cggcgcaata attcggtcta gtaacaaatt cattgaagaa tttgattcgc
                                                                      1740
cgatttgtgg tgtgacagtt gttcgaccgc ggatgacaga cggtgaggtt ttggagaaca
                                                                      1800
tcatgccgga agatgcacca tatcatgaca tttqcaagtt cattttqaqq ctcacatcaq
                                                                      1860
aaagtgtaac tttctcagga gaggggccag aagttagtga tttgaacgaa aaatggggaa
                                                                      1920
caattgtgta Ctatgagaaa aatttgcaaa ttggcgagaa aaaatgttcg agaggaaatt
                                                                      1980
tccacgtgga tggcggattc atttgctctg agaatcgtta cagtctcgga cttgagccaa
                                                                      2040
atccaattag agaaccagtg gcgtttaaag ttcgtaaagc aatagtggat ggaattcgct
                                                                      2100
tttcctacaa aaaagacggg agtgtttggc ttcaaaaccg catgaagtac ccggtatttg
                                                                      2160
teacttetgg gtatetegae gagcaateag gaggeetaaa gaaggataaa gtgcacaaag
                                                                      2220
tttacggatg tgcgtctatc aaaacgtttg gcttcaacgt ttccaaacaa atcatcagag
                                                                      2280
acgcgcttct ttccaagcaa atggcaacaa tgtacttgca aggaaaattg actccgatga
                                                                      2340
attatatcta cgagaagaag actcaggaag agctgcgaag ggaagcaaca cgcaccactg
                                                                      2400
attcattggc caagtactgt tgtgtccgtg tctcgttctg caaaggattt ggagaagcat
                                                                      2460
acccagaacg cccgtcaatt catgattgtc cagtttggat tgagttgaaa atcaacattg
                                                                      2520
cctacgattt catggattca atctgccagt acataaccaa ctgcttcgag ccgctaggaa
                                                                      2580
tggaagattt tgcaaaattg ggaatcaacg tcagtgatga ctaaatgata acttttttca
                                                                      2640
ctcaccctac tagatactga tttagtctta ttccaaatca tccaacqata tcaaactttt
                                                                      2700
tcctttgaac tttgcatact atgttatcac aagttccaag cagtttcaat acaaacatag
                                                                      2760
gatatgttaa caacttttga taagaatcaa gttaccaact gttcattgtg agctttgagc
                                                                      2820
tgtatagaag gacaatgtat cccatacctc aatctttaat agtcatcagt cactggtccc
                                                                      2880
gcaccaattt tttcgattcg catatgtcat atattgcacc gtggcccttt ttattgtaac
                                                                      2940
ttttaatata ttttcttccc aacttgtgaa tatgattgat gaaccaccat tttgagtaat
                                                                      3000
aaatgtattt tttgtgg
                                                                      3017
<210> 53
<211> 3119
<212> DNA
<213> Caenorhabditis elegans
<400> 53
gtaatcaaat tgtaaaggaa aaatattaat agtcagagta cacataaatg ggtgatcatc
                                                                        60
ataatttaac gggccttccc ggtacctcca tcccgccaca gttcaactat tctcaqcccq
                                                                       120
gtaccagcac cggaggcccg ctttatggtg gaaaaccttc tcatggattg gaagatattc
                                                                       180
ctgatgtaga ggaatatgag aggaacctgc tcggggctgg agcaggtttt aatctgctca
                                                                       240
atgtaggaaa tatggctaat gaatttaaac caataatcac attggacacg aaaccacctc
                                                                       300
gtgatgccaa caagtcattg gcattcaatg gcgggttgaa gctaatcact ccgaaaactg
                                                                       360
aagttcccga cgagcacaca ccgatgatgt caccagtgaa tacaactaca aagattctac
                                                                       420
aacggagtgg tattaaaatg gaaatcccgc catatttgga tccagacagt caggatgatg
                                                                       480
accoggaaga tggtgtcaac taccoggatc cagatttatt tgacacaaaa aacacaaata
                                                                       540
tgaccgagta cgatttggat gtgttgaagc ttggaaaacc agcagtagat gaagcacgga
                                                                       600
```

aaaagatcga agttcccgac gctagtgcgc cgccaaacaa aattgtagaa tatttgatgt

```
attatagaac gttaaaagaa agtgaactca tacaactgaa tgcgtatcgg acaaaacqaa
                                                                       720
atcgattatc gttgaacttg gtcaaaaaca atattgatcg agagttcgac caaaaagctt
                                                                       780
gcgagtccct ggtgaaaaaa ttgaaggata agaagaatga tctccagaac ctgattgatg
                                                                       840
tggttctttc aaaaggtaca aaatataccg gttgcattac aattccaagg acacttgatg
                                                                       900
gccggttaca ggtccacgga agaaaaggtt tccctcacgt agtctatggc aaactgtgga
                                                                       960
ggtttaatga aatgacaaaa aacgaaacgc gtcatgtgga ccactgcaag cacgcatttg
                                                                      1020
aaatgaaaag tgacatggta tgcgtgaatc cctatcacta cgaaattgtc attggaacta
                                                                      1080
tgattgttgg gcagagggat catgacaatc gagatatgcc gccgccacat caacqctacc
                                                                      1140
acactccagg tcggcaggat ccagttgacg atatgagtag atttatacca ccagcttcca
                                                                      1200
ttcgtccgcc tccgatgaac atgcacacaa ggcctcagcc tatgcctcaa caattgcctt
                                                                      1260
cagttggcgc aacgtttgcc catcetetee cacateagge gecacataae ceaggggttt
                                                                      1320
cacatccgta ctccattgct ccacagaccc attacccgtt gaacatgaac ccaattccgc
                                                                      1380
aaatgccgca aatgccacaa atgccaccac ctctccatca gggatatgga atgaatgggc
                                                                      1440
cgagttgctc ttcagaaaac aacaatccat tccaccaaaa tcaccattat aatgatatta
                                                                      1500
gccatccaaa tcactattcc tacgactgtg gtccgaactt gtacgggttt ccaactcctt
                                                                      1560
atcoggattt tcaccatcct ttcaatcagc aaccacaca gccgccacaa ctatcacaaa
                                                                      1620
accatacgtc ccaacaaggc agtcatcaac cagggcacca aggtcaggta ccgaatgatc
                                                                      1680
caccaatttc aagaccagtg ttacaaccat caacagtcac cttggacgtg ttccgtcggt
                                                                      1740
actgtagaca gacatttgga aatcgatttt ttgaaggaga aagtgaacaa tccggcgcaa
                                                                      1800
taattcggtc tagtaacaaa ttcattgaag aatttgattc gccgatttgt ggtgtgacag
                                                                      1860
ttgttcgacc gcggatgaca gacggtgagg ttttggagaa catcatgccg gaaqatgcac
                                                                      1920
catatcatga catttgcaag ttcattttga ggctcacatc agaaagtgta actttctcag
                                                                      1980
gagaggggcc agaagttagt gatttgaacg aaaaatgggg aacaattgtg tactatgaga
                                                                      2040
aaaatttgca aattggcgag aaaaaatgtt cgagaggaaa tttccacgtg gatggcggat
                                                                      2100
tcatttgctc tgagaatcgt tacagtctcg gacttgagcc aaatccaatt agagaaccag
                                                                      2160
tggcgtttaa agttcgtaaa gcaatagtgg atggaattcg cttttcctac aaaaaaqacq
                                                                      2220
ggagtgtttg gcttcaaaac cgcatgaagt acccggtatt tgtcacttct qqqtatctcq
                                                                      2280
acgagcaatc aggaggccta aagaaggata aagtgcacaa agtttacgga tgtgcgtcta
                                                                      2340
tcaaaacgtt tggcttcaac gtttccaaac aaatcatcag agacgcgctt ctttccaagc
                                                                      2400
aaatggcaac aatgtacttg caaggaaaat tgactccgat gaattatatc tacgagaaga
                                                                      2460
agactcagga agagctgcga agggaagcaa cacgcaccac tgattcattg gccaagtact
                                                                      2520
gttgtgtccg tgtctcgttc tgcaaaggat ttggagaagc atacccagaa cgcccgtcaa
                                                                      2580
ttcatgattg tccagtttgg attgagttga aaatcaacat tgcctacgat ttcatggatt
                                                                      2640
caatctgcca gtacataacc aactgcttcg agccgctagg aatggaagat tttgcaaaat
                                                                      2700
tgggaatcaa cgtcagtgat gactaaatga taactttttt cactcaccct actagatact
                                                                      2760
gatttagtct tattccaaat catccaacga tatcaaactt tttcctttga actttgcata
                                                                      2820
ctatgttatc acaagttcca agcagtttca atacaaacat aggatatgtt aacaactttt
                                                                      2880
gataagaatc aagttaccaa ctgttcattg tgagctttga gctgtataga aggacaatgt
                                                                      2940
atcccatacc tcaatcttta atagtcatca gtcactggtc ccgcaccaat tttttcgatt
                                                                      3000
cgcatatgtc atatattgca ccgtggccct ttttattgta acttttaata tattttcttc
                                                                      3060
ccaacttgtg aatatgattg atgaaccacc attttgagta ataaatgtat tttttgtgg
                                                                      3119
```

```
<210> 54
<211> 103
<212> PRT
```

<213> Caenorhabditis elegans

<400> 54

Lys Lys Thr Thr Arg Arg Asn Ala Trp Gly Asn Met Ser Tyr Ala 1 5 10 15 Glu Leu Ile Thr Thr Ala Ile Met Ala Ser Pro Glu Lys Arg Leu Thr 20 25 Leu Ala Gln Val Tyr Glu Trp Met Val Gln Asn Val Pro Tyr Phe Arg 35 40 Asp Lys Gly Asp Ser Asn Ser Ser Ala Gly Trp Lys Asn Ser Ile Arg 50 55 60 His Asn Leu Ser Leu His Ser Arg Phe Met Arg Ile Gln Asn Glu Gly 65 70 75 Ala Gly Lys Ser Ser Trp Trp Val Ile Asn Pro Asp Ala Lys Pro Gly

```
85 90 95
Met Asn Pro Arg Thr Arg
```

<210> 55 <211> 41

<212> PRT

<213> Caenorhabditis elegans

<400> 55

Thr Phe Met Asn Thr Pro Asp Asp Val Met Met Asn Asp Asp Met Glu

1 10 15

Pro Ile Pro Arg Asp Arg Cys Asn Thr Trp Pro Met Arg Arg Pro Gln 20 25 30

Leu Glu Pro Pro Leu Asn Ser Ser Pro 35 40

<210> 56

<211> 109

<212> PRT

<213> Caenorhabditis elegans

<400> 56

Asp Asp Thr Val Ser Gly Lys Lys Thr Thr Thr Arg Arg Asn Ala Trp $1 \hspace{1cm} 5 \hspace{1cm} 10 \hspace{1cm} 15$

Gly Asn Met Ser Tyr Ala Glu Leu Ile Thr Thr Ala Ile Met Ala Ser 20 25 30

Pro Glu Lys Arg Leu Thr Leu Ala Gln Val Tyr Glu Trp Met Val Gln 35 40 45

Asn Val Pro Tyr Phe Arg Asp Lys Gly Asp Ser Asn Ser Ser Ala Gly 50 55

Trp Lys Asn Ser Ile Arg His Asn Leu Ser Leu His Ser Arg Phe Met 65 75 80

Arg Ile Gln Asn Glu Gly Ala Gly Lys Ser Ser Trp Trp Val Ile Asn 85 90 95

Pro Asp Ala Lys Pro Gly Met Asn Pro Arg Arg Thr Arg 100 105

<210> 57

<211> 655

<212> PRT

<213> Homo sapiens

<400> 57

Met Ala Glu Ala Pro Gln Val Val Glu Ile Asp Pro Asp Phe Glu Pro
1 5 10 15

Leu Pro Arg Pro Arg Ser Cys Thr Trp Pro Leu Pro Arg Pro Glu Phe 20 25 30

Ser Gln Ser Asn Ser Ala Thr Ser Ser Pro Ala Pro Ser Gly Ser Ala 35 40 45

Ala Ala Asn Pro Asp Ala Ala Ala Gly Leu Pro Ser Ala Ser Ala Ala 50 55 60

Ala Val Ser Ala Asp Phe Met Ser Asn Leu Ser Leu Leu Glu Glu Ser 65 70 75 80

Glu Asp Phe Pro Gln Ala Pro Gly Ser Val Ala Ala Ala Val Ala Ala

```
Ala Ala Ala Ala Ala Thr Gly Gly Leu Cys Gly Asp Phe Gln Gly
            100
                                105
Pro Glu Ala Gly Cys Leu His Pro Ala Pro Pro Gln Pro Pro Pro Pro
        115
                            120
                                                125
Gly Pro Val Ser Gln His Pro Pro Val Pro Pro Ala Ala Ala Gly Pro
                        135
                                            140
Leu Ala Gly Gln Pro Arg Lys Ser Ser Ser Ser Arg Arg Asn Ala Trp
                    150
                                        155
Gly Asn Leu Ser Tyr Ala Asp Leu Ile Thr Lys Ala Ile Glu Ser Ser
                165
                                    170
                                                        175
Ala Glu Lys Arg Leu Thr Leu Ser Gln Ile Tyr Glu Trp Met Val Lys
                                185
            180
Ser Val Pro Tyr Phe Lys Asp Lys Gly Asp Ser Asn Ser Ser Ala Gly
                            200
        195
                                                205
Trp Lys Asn Ser Ile Arg His Asn Leu Ser Leu His Ser Lys Phe Ile
    210
                        215
                                            220
Arg Val Gln Asn Glu Gly Thr Gly Lys Ser Ser Trp Trp Met Leu Asn
225
                   230
                                       235
Pro Glu Gly Gly Lys Ser Gly Lys Ser Pro Arg Arg Ala Ala Ser
                245
                                   250
Met Asp Asn Asn Ser Lys Phe Ala Lys Ser Arg Ser Arg Ala Ala Lys
           260
                                265
                                                    270
Lys Lys Ala Ser Leu Gln Ser Gly Gln Glu Gly Ala Gly Asp Ser Pro
                            280
                                                285
Gly Ser Gln Phe Ser Lys Trp Pro Ala Ser Pro Gly Ser His Ser Asn
                                            300
                       295
Asp Asp Phe Asp Asn Trp Ser Thr Phe Arg Pro Arg Thr Ser Ser Asn
305
                                       315
                   310
Ala Ser Thr Ile Ser Gly Arg Leu Ser Pro Ile Met Thr Glu Gln Asp
                325
                                    330
Asp Leu Gly Glu Gly Asp Val His Ser Met Val Tyr Pro Pro Ser Ala
            340
                                345
                                                    350
Ala Lys Met Ala Ser Thr Leu Pro Ser Leu Ser Glu Ile Ser Asn Pro
       355
                            360
                                                365
Glu Asn Met Glu Asn Leu Leu Asp Asn Leu Asn Leu Leu Ser Ser Pro
                        375
                                            380
Thr Ser Leu Thr Val Ser Thr Gln Ser Ser Pro Gly Thr Met Met Gln
                    390
                                        395
Gln Thr Pro Cys Tyr Ser Phe Ala Pro Pro Asn Thr Ser Leu Asn Ser
                405
                                    410
Pro Ser Pro Asn Tyr Gln Lys Tyr Thr Tyr Gly Gln Ser Ser Met Ser
            420
                                425
Pro Leu Pro Gln Met Pro Ile Gln Thr Leu Gln Asp Asn Lys Ser Ser
        435
                            440
                                                445
Tyr Gly Gly Met Ser Gln Tyr Asn Cys Ala Pro Gly Leu Leu Lys Glu
    450
                        455
                                            460
Leu Leu Thr Ser Asp Ser Pro Pro His Asn Asp Ile Met Thr Pro Val
                   470
                                        475
Asp Pro Gly Val Ala Gln Pro Asn Ser Arg Val Leu Gly Gln Asn Val
                485
                                    490
Met Met Gly Pro Asn Ser Val Met Ser Thr Tyr Gly Ser Gln Ala Ser
            500
                                505
                                                    510
His Asn Lys Met Met Asn Pro Ser Ser His Thr His Pro Gly His Ala
       515
                            520
                                                525
Gln Gln Thr Ser Ala Val Asn Gly Arg Pro Leu Pro His Thr Val Ser
                       535
                                            540
Thr Met Pro His Thr Ser Gly Met Asn Arg Leu Thr Gln Val Lys Thr
                    550
                                        555
Pro Val Gln Val Pro Leu Pro His Pro Met Gln Met Ser Ala Leu Gly
```

```
565
                                    570
Gly Tyr Ser Ser Val Ser Ser Cys Asn Gly Tyr Gly Arg Met Gly Leu
                                585
Leu His Gln Glu Lys Leu Pro Ser Asp Leu Asp Gly Met Phe Ile Glu
                            600
Arg Leu Asp Cys Asp Met Glu Ser Ile Ile Arg Asn Asp Leu Met Asp
                        615
                                             620
Gly Asp Thr Leu Asp Phe Asn Phe Asp Asn Val Leu Pro Asn Gln Ser
                    630
                                         635
Phe Pro His Ser Val Lys Thr Thr His Ser Trp Val Ser Gly
<210> 58
<211> 98
<212> PRT
<213> Caenorhabditis elegans
<400> 58
Lys Pro Asn Pro Trp Gly Glu Glu Ser Tyr Ser Asp Ile Ile Ala Lys
                                    10
Ala Leu Glu Ser Ala Pro Asp Gly Arg Leu Lys Leu Asn Glu Ile Tyr
            20
                                25
Gln Trp Phe Ser Asp Asn Ile Pro Tyr Phe Gly Glu Arg Ser Ser Pro
                            40
Glu Glu Ala Ala Gly Trp Lys Asn Ser Ile Arg His Asn Leu Ser Leu
                        5.5
                                             60
His Ser Arg Phe Met Arg Ile Gln Asn Glu Gly Ala Gly Lys Ser Ser
                    70
                                        75
Trp Trp Val Ile Asn Pro Asp Ala Lys Pro Gly Met Asn Pro Arg Arg
Thr Arg
<210> 59
<211> 7
<212> PRT
<213> Caenorhabditis elegans
<400> 59
Trp Lys Asn Ser Ile Arg His
<210> 60
<211> 121
<212> PRT
<213> Caenorhabditis elegans
<400> 60
Gln Val Leu Asp Asp His Asp Tyr Gly Arg Cys Val Asp Trp Trp Gly
Val Gly Val Val Met Tyr Glu Met Met Cys Gly Arg Leu Pro Phe Tyr
            20
                                25
Ser Lys Asp His Asn Lys Leu Phe Glu Leu Ile Met Ala Gly Asp Leu
                            40
Arg Phe Pro Ser Lys Leu Ser Gln Glu Ala Arg Thr Leu Leu Thr Gly
```

```
Leu Leu Val Lys Asp Pro Thr Gln Arg Leu Gly Gly Pro Glu Asp
                    70
                                        75
65
                                                             80
Ala Leu Glu Ile Cys Arg Ala Asp Phe Phe Arg Thr Val Asp Trp Glu
                                    90
                                                         95
                85
Ala Thr Tyr Arg Lys Glu Ile Glu Pro Pro Tyr Lys Pro Asn Val Gln
            100
                                105
Ser Glu Thr Asp Thr Ser Tyr Phe Asp
        115
                            120
<210> 61
<211> 66
<212> PRT
<213> Caenorhabditis elegans
<400> 61
Thr Met Glu Asp Phe Asp Phe Leu Lys Val Leu Gly Lys Gly Thr Phe
Gly Lys Val Ile Leu Cys Lys Glu Lys Arg Thr Gln Lys Leu Tyr Ala
                                25
Ile Lys Ile Leu Lys Lys Asp Val Ile Ile Ala Arg Glu Glu Val Ala
                            40
His Thr Leu Thr Glu Asn Arg Val Leu Gln Arg Cys Lys His Pro Phe
                        55
                                            60
Leu Thr
65
<210> 62
<211> 45
<212> PRT
<213> Caenorhabditis elegans
<400> 62
Lys Leu Glu Asn Leu Leu Asp Lys Asp Gly His Ile Lys Ile Ala
1
                                    10
Asp Phe Gly Leu Cys Lys Glu Glu Ile Ser Phe Gly Asp Lys Thr Ser
                                25
Thr Phe Cys Gly Thr Pro Glu Tyr Leu Ala Pro Glu Val
                            40
<210> 63
<211> 57
<212> PRT
<213> Caenorhabditis elegans
<400> 63
Tyr Phe Gln Glu Leu Lys Tyr Ser Phe Gln Glu Gln His Tyr Leu Cys
1
                                    10
Phe Val Met Gln Phe Ala Asn Gly Gly Glu Leu Phe Thr His Val Arg
                                25
Lys Cys Gly Thr Phe Ser Glu Pro Arg Ala Arg Phe Tyr Gly Ala Glu
```

Ile Val Leu Ala Leu Gly Tyr Leu His

```
<211> 59
<212> PRT
<213> Caenorhabditis elegans
<400> 64
Ser Thr Phe Ala Ile Phe Tyr Phe Gln Thr Met Leu Phe Glu Lys Pro
Arg Pro Asn Met Phe Met Val Arg Cys Leu Gln Trp Thr Thr Val Ile
                                25
Glu Arg Thr Phe Tyr Ala Glu Ser Ala Glu Val Arg Gln Arg Trp Ile
His Ala Ile Glu Ser Ile Ser Lys Lys Tyr Lys
<210> 65
<211> 33
<212> PRT
<213> Caenorhabditis elegans
<400> 65
Leu Gln Glu Leu Lys Tyr Ser Phe Gln Thr Asn Asp Arg Leu Cys Phe
Val Met Glu Phe Ala Ile Gly Gly Asp Leu Tyr Tyr His Leu Asn Arg
                                25
Glu
<210> 66
<211> 21
<212> PRT
<213> Caenorhabditis elegans
<400> 66
Val Val Ile Glu Gly Trp Leu His Lys Lys Gly Glu His Ile Arg Asn
Trp Arg Pro Arg Phe
            20
<210> 67
<211> 26
<212> PRT
<213> Caenorhabditis elegans
<400> 67
Phe Ser Glu Pro Arg Ala Arg Phe Tyr Gly Ser Glu Ile Val Leu Ala
                                    10
Leu Gly Tyr Leu His Ala Asn Ser Ile Val
            20
<210> 68
<211> 39
<212> PRT
<213> Caenorhabditis elegans
```

<400> 68

```
Ile Arg Val Ser Phe Cys Lys Gly Phe Gly Glu Thr Tyr Ser Arg Leu
                 5
                                     10
Lys Val Val Asn Leu Pro Cys Trp Ile Glu Ile Ile Leu His Glu Pro
                                 25
Ala Asp Glu Tyr Asp Thr Val
        35
<210> 69
<211> 45
<212> PRT
<213> Caenorhabditis elegans
<400> 69
Ser Arg Asn Ser Lys Ser Ser Gln Ile Arg Asn Thr Val Gly Ala Gly
                                     10
Ile Gln Leu Ala Tyr Glu Asn Gly Glu Leu Trp Leu Thr Val Leu Thr
            20
                                25
Asp Gln Ile Val Phe Val Gln Cys Pro Phe Leu Asn Gln
                            40
<210> 70
<211> 29
<212> PRT
<213> Caenorhabditis elegans
<400> 70
Asn Glu Met Leu Asp Pro Glu Pro Lys Tyr Pro Lys Glu Glu Lys Pro
1
                                     10
Trp Cys Thr Ile Phe Tyr Tyr Glu Leu Thr Val Arg Val
            20
<210> 71
<211> 29
<212> PRT
<213> Caenorhabditis elegans
<400> 71
Gln Leu Gly Lys Ala Phe Glu Ala Lys Val Pro Thr Ile Thr Ile Asp
                                     10
Gly Ala Thr Gly Ala Ser Asp Glu Cys Arg Met Ser Leu
            20
                                25
<210> 72
<211> 105
<212> PRT
<213> Caenorhabditis elegans
<400> 72
Ser Pro Asp Asp Gly Leu Leu Asp Ser Ser Glu Glu Ser Arg Arg
1
                                                         15
Gln Lys Thr Cys Arg Val Cys Gly Asp His Ala Thr Gly Tyr Asn Phe
            20
                                25
Asn Val Ile Thr Cys Glu Ser Cys Lys Ala Phe Phe Arg Arg Asn Ala
                            40
```

Leu Arg Pro Lys Glu Phe Lys Cys Pro Tyr Ser Glu Asp Cys Glu Ile

```
Asn Ser Val Ser Arg Arg Phe Cys Gln Lys Cys Arg Leu Arg Lys Cys
                    70
Phe Thr Val Gly Met Lys Lys Glu Trp Ile Leu Asn Glu Glu Gln Leu
                                     90
Arg Arg Arg Lys Asn Ser Arg Leu Asn
            100
<210> 73
<211> 89
<212> PRT
<213> Caenorhabditis elegans
<400> 73
Leu Asp Ser Ser Glu Glu Ser Arg Arg Arg Gln Lys Thr Cys Arg Val
Cys Gly Asp His Ala Thr Gly Tyr Asn Phe Asn Val Ile Thr Cys Glu
                                25
Ser Cys Lys Ala Phe Phe Arg Arg Asn Ala Leu Arg Pro Lys Glu Phe
                                                 45
Lys Cys Pro Tyr Ser Glu Asp Cys Glu Ile Asn Ser Val Ser Arg Arg
                        55
Phe Cys Gln Lys Cys Arg Leu Arg Lys Cys Phe Thr Val Gly Met Lys
Lys Glu Trp Ile Leu Asn Glu Glu Gln
<210> 74
<211> 73
<212> PRT
<213> Caenorhabditis elegans
<400> 74
Asp Ile Met Asn Ile Met Asp Val Thr Met Arg Arg Phe Val Lys Val
Ala Lys Gly Val Pro Ala Phe Arg Glu Val Ser Gln Glu Gly Lys Phe
Ser Leu Leu Lys Gly Gly Met Ile Glu Met Leu Thr Val Arg Gly Val
                            40
Thr Arg Tyr Asp Ala Ser Thr Asn Ser Phe Lys Thr Pro Thr Ile Lys
Gly Gln Asn Val Ser Val Asn Val Asp
<210> 75
<211> 112
<212> PRT
<213> Caenorhabditis elegans
Ser Gly Ser Leu Val Asp Leu Met Ile Lys Asn Leu Thr Ala Tyr Thr
                                    10
Gln Gly Leu Asn Glu Thr Val Lys Asn Arg Thr Ala Glu Leu Glu Lys
                                2.5
Glu Gln Glu Lys Gly Asp Gln Leu Leu Met Glu Leu Leu Pro Lys Ser
```

Val Ala Asn Asp Leu Lys Asn Gly Ile Ala Val Asp Pro Lys Val Tyr 55 Glu Asn Ala Thr Ile Leu Tyr Ser Asp Ile Val Gly Phe Thr Ser Leu 70 75 80 Cys Ser Gln Ser Gln Pro Met Glu Val Val Thr Leu Leu Ser Gly Met 90 95 Tyr Gln Arg Phe Asp Leu Ile Ile Ser Gln Gln Gly Gly Tyr Lys Val 105 110 <210> 76 <211> 107 <212> PRT <213> Caenorhabditis elegans <400> 76 Met Glu Thr Ile Gly Asp Ala Tyr Cys Val Ala Ala Gly Leu Pro Val Val Met Glu Lys Asp His Val Lys Ser Ile Cys Met Ile Ala Leu Leu 25 Gln Arg Asp Cys Leu His His Phe Glu Ile Pro His Arg Pro Gly Thr Phe Leu Asn Cys Arg Trp Gly Phe Asn Ser Gly Pro Val Phe Ala Gly 55 Val Ile Gly Gln Lys Ala Pro Arg Tyr Ala Cys Phe Gly Glu Ala Val 75 Ile Leu Ala Ser Lys Met Glu Ser Ser Gly Val Glu Asp Arg Ile Gln 90 Met Thr Leu Ala Ser Gln Gln Leu Leu Glu Glu <210> 77 <211> 43 <212> PRT <213> Caenorhabditis elegans <400> 77 Asp Ile Leu Lys Gly Leu Glu Tyr Ile His Ala Ser Ala Ile Asp Phe 1 10 15 His Gly Asn Leu Thr Leu His Asn Cys Met Leu Asp Ser His Trp Ile 25 Val Lys Leu Ser Gly Phe Gly Val Asn Arg Leu <210> 78 <211> 15 <212> PRT <213> Caenorhabditis elegans <400> 78 Asp Met Tyr Ser Phe Gly Val Ile Leu His Glu Ile Ile Leu Lys <210> 79

<211> 67 <212> PRT

<213> Caenorhabditis elegans <400> 79 Ala Ile Lys Ile Asn Val Asp Asp Pro Ala Ser Thr Glu Asn Leu Asn 1 Tyr Leu Met Glu Ala Asn Ile Met Lys Asn Phe Lys Thr Asn Phe Ile 25 Val Gln Leu Tyr Gly Val Ile Ser Thr Val Gln Pro Ala Met Val Val Met Glu Met Met Asp Leu Gly Asn Leu Arg Asp Tyr Leu Arg Ser Lys 55 60 Arg Glu Asp 65 <210> 80 <211> 54 <212> PRT <213> Caenorhabditis elegans <400> 80 Val Ile Lys Lys Pro Glu Cys Cys Glu Asn Tyr Trp Tyr Lys Val Met Lys Met Cys Trp Arg Tyr Ser Pro Arg Asp Arg Pro Thr Phe Leu Gln 25 Leu Val His Leu Leu Ala Ala Glu Ala Ser Pro Glu Phe Arg Asp Leu 40 Ser Phe Val Leu Thr Asp 50 <210> 81 <211> 69 <212> PRT <213> Caenorhabditis elegans <400> 81 Lys Gln Asp Ser Gly Met Ala Ser Glu Leu Lys Asp Ile Phe Ala Asn Ile His Thr Ile Thr Gly Tyr Leu Leu Val Arg Gln Ser Ser Pro Phe Ile Ser Leu Asn Met Phe Arg Asn Leu Arg Arg Ile Glu Ala Lys Ser 40 Leu Phe Arg Asn Leu Tyr Ala Ile Thr Val Phe Glu Asn Pro Asn Leu Lys Lys Leu Phe Asp 65 <210> 82 <211> 52 <212> PRT <213> Caenorhabditis elegans <400> 82 Phe Pro His Leu Arg Glu Ile Thr Gly Thr Leu Leu Val Phe Glu Thr

10

Glu Gly Leu Val Asp Leu Arg Lys Ile Phe Pro Asn Leu Arg Val Ile

25

```
Gly Gly Arg Ser Leu Ile Gln His Tyr Ala Leu Ile Ile Tyr Arg Asn
        35
Pro Asp Leu Glu
    50
<210> 83
<211> 46
<212> PRT
<213> Caenorhabditis elegans
<400> 83
Glu Ile Gly Leu Asp Lys Leu Ser Val Ile Arg Asn Gly Gly Val Arg
1
                                    10
Ile Ile Asp Asn Arg Lys Leu Cys Tyr Thr Lys Thr Ile Asp Trp Lys
            20
                                25
His Leu Ile Thr Ser Ser Ile Asn Asp Val Val Asp Asn
        35
                            40
<210> 84
<211> 36
<212> PRT
<213> Caenorhabditis elegans
<400> 84
Tyr Asn Ala Asp Asp Trp Glu Leu Arg Gln Asp Asp Val Val Leu Gly
1
                 5
Gln Gln Cys Gly Glu Gly Ser Phe Gly Lys Val Tyr Leu Gly Thr Gly
            20
                                25
                                                     30
Asn Asn Val Val
        35
<210> 85
<211> 24
<212> PRT
<213> Caenorhabditis elegans
<400> 85
Asp Ser Leu Ala Lys Tyr Cys Cys Val Arg Val Ser Phe Cys Lys Gly
                                                         15
Phe Gly Glu Ala Tyr Pro Glu Arg
            20
<210> 86
<211> 13
<212> PRT
<213> Caenorhabditis elegans
<400> 86
Gly Trp Asp Trp Ile Val Ala Pro Pro Arg Tyr Asn Ala
                 5
<210> 87
<211> 121
<212> PRT
```

<213> Homo sapiens

```
<400> 87
Glu Val Leu Glu Asp Asn Asp Tyr Gly Arg Ala Val Asp Trp Trp Gly
Leu Gly Val Val Met Tyr Glu Met Met Cys Gly Arg Leu Pro Phe Tyr
Asn Gln Asp His Glu Lys Leu Phe Glu Leu Ile Leu Met Glu Glu Ile
Arg Phe Pro Arg Thr Leu Gly Pro Glu Ala Lys Ser Leu Leu Ser Gly
Leu Leu Lys Lys Asp Pro Thr Gln Arg Leu Gly Gly Ser Glu Asp
Ala Lys Glu Ile Met Gln His Arg Phe Phe Ala Asn Ile Val Trp Gln
Asp Val Tyr Glu Lys Lys Leu Ser Pro Pro Phe Lys Pro Gln Val Thr
                                105
Ser Glu Thr Asp Thr Arg Tyr Phe Asp
<210> 88
<211> 121
<212> PRT
<213> Caenorhabditis elegans
<400> 88
Gln Val Leu Asp Asp His Asp Tyr Gly Arg Cys Val Asp Trp Trp Gly
Val Gly Val Val Met Tyr Glu Met Met Cys Gly Arg Leu Pro Phe Tyr
Ser Lys Asp His Asn Lys Leu Phe Glu Leu Ile Met Ala Gly Asp Leu
                            40
Arg Phe Pro Ser Lys Leu Ser Gln Glu Ala Arg Thr Leu Leu Thr Gly
                        55
Leu Leu Val Lys Asp Pro Thr Gln Arg Leu Gly Gly Pro Glu Asp
Ala Leu Glu Ile Cys Arg Ala Asp Phe Phe Arg Thr Val Asp Trp Glu
Ala Thr Tyr Arg Lys Glu Ile Glu Pro Pro Tyr Lys Pro Asn Val Gln
            100
Ser Glu Thr Asp Thr Ser Tyr Phe Asp
<210> 89
<211> 66
<212> PRT
<213> Homo sapiens
<400> 89
Thr Met Asn Glu Phe Glu Tyr Leu Lys Leu Leu Gly Lys Gly Thr Phe
                                    10
Gly Lys Val Ile Leu Val Lys Glu Lys Ala Thr Gly Arg Tyr Tyr Ala
Met Lys Ile Leu Lys Lys Glu Val Ile Val Ala Lys Asp Glu Val Ala
```

35 40 45 His Thr Leu Thr Glu Asn Arg Val Leu Gln Asn Ser Arg His Pro Phe

```
Leu Thr
65
<210> 90
<211> 66
<212> PRT
<213> Caenorhabditis elegans
<400> 90
Thr Met Glu Asp Phe Asp Phe Leu Lys Val Leu Gly Lys Gly Thr Phe
Gly Lys Val Ile Leu Cys Lys Glu Lys Arg Thr Gln Lys Leu Tyr Ala
                                25
Ile Lys Ile Leu Lys Lys Asp Val Ile Ile Ala Arg Glu Glu Val Ala
                            40
His Thr Leu Thr Glu Asn Arg Val Leu Gln Arg Cys Lys His Pro Phe
                        55
                                             60
Leu Thr
65
<210> 91
<211> 45
<212> PRT
<213> Homo sapiens
<400> 91
Lys Leu Glu Asn Leu Met Leu Asp Lys Asp Gly His Ile Lys Ile Thr
                                    10
Asp Phe Gly Leu Cys Lys Glu Gly Ile Lys Asp Gly Ala Thr Met Lys
Thr Phe Cys Gly Thr Pro Glu Tyr Leu Ala Pro Glu Val
        35
                            40
<210> 92
<211> 45
<212> PRT
<213> Caenorhabditis elegans
<400> 92
Lys Leu Glu Asn Leu Leu Asp Lys Asp Gly His Ile Lys Ile Ala
                                    10
Asp Phe Gly Leu Cys Lys Glu Glu Ile Ser Phe Gly Asp Lys Thr Ser
                                25
Thr Phe Cys Gly Thr Pro Glu Tyr Leu Ala Pro Glu Val
                            40
<210> 93
<211> 57
<212> PRT
<213> Homo sapiens
<400> 93
Phe Leu Thr Ala Leu Lys Tyr Ser Phe Gln Thr His Asp Arg Leu Cys
1
                                    10
Phe Val Met Glu Tyr Ala Asn Gly Gly Glu Leu Phe Phe His Leu Ser
```

```
25
Arg Glu Arg Val Phe Ser Glu Asp Arg Ala Arg Phe Tyr Gly Ala Glu
                            40
                                                 45
Ile Val Ser Ala Leu Asp Tyr Leu His
                        55
<210> 94
<211> 57
<212> PRT
<213> Caenorhabditis elegans
<400> 94
Tyr Phe Gln Glu Leu Lys Tyr Ser Phe Gln Glu Gln His Tyr Leu Cys
1
Phe Val Met Gln Phe Ala Asn Gly Gly Glu Leu Phe Thr His Val Arg
                                25
                                                     30
Lys Cys Gly Thr Phe Ser Glu Pro Arg Ala Arg Phe Tyr Gly Ala Glu
                            40
                                                 45
Ile Val Leu Ala Leu Gly Tyr Leu His
                        55
<210> 95
<211> 59
<212> PRT
<213> Homo sapiens
<400> 95
Asn Asn Phe Ser Val Ala Gln Cys Gln Leu Met Lys Thr Glu Arg Pro
                5
                                    10
Arg Pro Asn Thr Phe Ile Ile Arg Cys Leu Gln Trp Thr Thr Val Ile
            20
                                25
Glu Arg Thr Phe His Val Glu Thr Pro Glu Glu Arg Glu Glu Trp Ala
                            40
Thr Ala Ile Gln Thr Val Ala Asp Gly Leu Lys
   50
                        55
<210> 96
<211> 59
<212> PRT
<213> Caenorhabditis elegans
<400> 96
Ser Thr Phe Ala Ile Phe Tyr Phe Gln Thr Met Leu Phe Glu Lys Pro
                                    10
                                                         15
Arg Pro Asn Met Phe Met Val Arg Cys Leu Gln Trp Thr Thr Val Ile
            20
                                25
Glu Arg Thr Phe Tyr Ala Glu Ser Ala Glu Val Arg Gln Arg Trp Ile
                            40
His Ala Ile Glu Ser Ile Ser Lys Lys Tyr Lys
   50
                        55
<210> 97
<211> 33
<212> PRT
<213> Homo sapiens
```

```
<400> 97
Leu Thr Ala Leu Lys Tyr Ser Phe Gln Thr His Asp Arg Leu Cys Phe
                                     10
Val Met Glu Tyr Ala Asn Gly Gly Glu Leu Phe Phe His Leu Ser Arg
                                 25
Glu
<210> 98
<211> 33
<212> PRT
<213> Caenorhabditis elegans
<400> 98
Leu Gln Glu Leu Lys Tyr Ser Phe Gln Thr Asn Asp Arg Leu Cys Phe
                                    10
Val Met Glu Phe Ala Ile Gly Gly Asp Leu Tyr Tyr His Leu Asn Arg
                                25
Glu
<210> 99
<211> 36
<212> PRT
<213> Homo sapiens or Caenorhabditis elegans
<400> 99
Lys Leu Glu Asn Leu Leu Asp Lys Asp Gly His Ile Lys Ile Asp Phe
                                    10
Gly Leu Cys Lys Glu Ile Gly Thr Phe Cys Gly Thr Pro Glu Tyr Leu
            20
Ala Pro Glu Val
        35
<210> 100
<211> 37
<212> PRT
<213> Homo sapiens or Caenorhabditis elegans
<400> 100
Leu Lys Tyr Ser Phe Gln Leu Cys Phe Val Met Ala Asn Gly Glu Glu
                                    10
Leu Phe His Phe Ser Glu Arg Ala Arg Phe Tyr Gly Ala Glu Ile Val
            20
                                25
Ala Leu Tyr Leu His
        35
<210> 101
<211> 29
<212> PRT
<213> Homo sapiens or Caenorhabditis elegans
<400> 101
Phe Gln Met Glu Pro Arg Pro Asn Phe Arg Cys Leu Gln Trp Thr Thr
```

```
Val Ile Glu Arg Thr Phe Glu Glu Arg Trp Ala Ile Lys
            20
<210> 102
<211> 24
<212> PRT
<213> Homo sapiens or Caenorhabditis elegans
<400> 102
Leu Leu Lys Tyr Ser Phe Gln Thr Asp Arg Leu Cys Phe Val Met Glu
                                    10
Ala Gly Gly Leu His Leu Arg Glu
            20
<210> 103
<211> 366
<212> PRT
<213> Homo sapiens
<400> 103
Arg Gly Ala Ile Arg Ile Glu Lys Asn Ala Asp Leu Cys Tyr Leu Ser
1
                                    10
Thr Val Asp Trp Ser Leu Ile Leu Asp Ala Val Ser Asn Asn Tyr Ile
                                25
Val Gly Asn Lys Pro Pro Lys Glu Cys Gly Asp Leu Cys Pro Gly Thr
Met Glu Glu Lys Pro Met Cys Glu Lys Thr Thr Ile Asn Asn Glu Tyr
                        55
Asn Tyr Arg Cys Trp Thr Thr Asn Arg Cys Gln Lys Met Cys Pro Ser
Thr Cys Gly Lys Arg Ala Cys Thr Glu Asn Asn Glu Cys Cys His Pro
Glu Cys Leu Gly Ser Cys Ser Ala Pro Asp Asn Asp Thr Ala Cys Val
            100
                                105
Ala Cys Arg His Tyr Tyr Ala Gly Val Cys Val Pro Ala Cys Pro
        115
                            120
                                                 125
Pro Asn Thr Tyr Arg Phe Glu Gly Trp Arg Cys Val Asp Arg Asp Phe
                        135
                                            140
Cys Ala Asn Ile Leu Ser Ala Glu Ser Ser Asp Ser Glu Gly Phe Val
                    150
                                        155
Ile His Asp Gly Glu Cys Met Gln Glu Cys Pro Ser Gly Phe Ile Arg
                165
                                    170
Asn Gly Ser Gln Ser Met Tyr Cys Ile Pro Cys Glu Gly Pro Cys Pro
            180
                                185
                                                     190
Lys Val Cys Glu Glu Glu Lys Lys Thr Lys Thr Ile Asp Ser Val Thr
        195
                            200
                                                 205
Ser Ala Gln Met Leu Gln Gly Cys Thr Ile Phe Lys Gly Asn Leu Leu
                        215
                                            220
Ile Asn Ile Arg Arg Gly Asn Asn Ile Ala Ser Glu Leu Glu Asn Phe
                    230
                                        235
Met Gly Leu Ile Glu Val Val Thr Gly Tyr Val Lys Ile Arg His Ser
```

His Ala Leu Val Ser Leu Ser Phe Leu Lys Asn Leu Arg Leu Ile Leu 265

Gly Glu Glu Gln Leu Glu Gly Asn Tyr Ser Phe Tyr Val Leu Asp Asn 280 Gln Asn Leu Gln Gln Leu Trp Asp Trp Asp His Arg Asn Leu Thr Ile

255

270

```
290
                        295
Lys Ala Gly Lys Met Tyr Phe Ala Phe Asn Pro Lys Leu Cys Val Ser
                    310
                                        315
Glu Ile Tyr Arg Met Glu Glu Val Thr Gly Thr Lys Gly Arg Gln Ser
                325
                                    330
Lys Gly Asp Ile Asn Thr Arg Asn Asn Gly Glu Arg Ala Ser Cys Glu
                                345
Ser Asp Val Leu His Phe Thr Ser Thr Thr Thr Ser Lys Asn
                            360
<210> 104
<211> 370
<212> PRT
<213> Homo sapiens
<400> 104
Arg Gly Ser Val Arg Ile Glu Lys Asn Asn Glu Leu Cys Tyr Leu Ala
Thr Ile Asp Trp Ser Arg Ile Leu Asp Ser Val Glu Asp Asn Tyr Ile
                                25
Val Leu Asn Lys Asp Asp Asn Glu Glu Cys Gly Asp Ile Cys Pro Gly
                            40
Thr Ala Lys Gly Lys Thr Asn Cys Pro Ala Thr Val Ile Asn Gly Gln
                        55
Phe Val Glu Arg Cys Trp Thr His Ser His Cys Gln Lys Val Cys Pro
                   70
                                        75
Thr Ile Cys Lys Ser His Gly Cys Thr Ala Glu Gly Leu Cys Cys His
                                    90
                85
Ser Glu Cys Leu Gly Asn Cys Ser Gln Pro Asp Asp Pro Thr Lys Cys
                                105
                                                    110
Val Ala Cys Arg Asn Phe Tyr Leu Asp Gly Arg Cys Val Glu Thr Cys
                           120
                                                125
Pro Pro Pro Tyr Tyr His Phe Gln Asp Trp Arg Cys Val Asn Phe Ser
                        135
                                            140
Phe Cys Gln Asp Leu His His Lys Cys Lys Asn Ser Arg Arg Gln Gly
                   150
                                        155
Cys His Gln Tyr Val Ile His Asn Asn Lys Cys Ile Pro Glu Cys Pro
                                   170
                                                        175
                165
Ser Gly Tyr Thr Met Asn Ser Ser Asn Leu Leu Cys Thr Pro Cys Leu
           180
                                185
                                                    190
Gly Pro Cys Pro Lys Val Cys His Leu Leu Glu Gly Glu Lys Thr Ile
                            200
       195
                                                205
Asp Ser Val Thr Ser Ala Gln Glu Leu Arg Gly Cys Thr Val Ile Asn
                        215
                                            220
Gly Ser Leu Ile Ile Asn Ile Arg Gly Gly Asn Asn Leu Ala Ala Glu
                   230
                                        235
Leu Glu Ala Asn Leu Gly Leu Ile Glu Glu Ile Ser Gly Tyr Leu Lys
                245
                                    250
Ile Arg Arg Ser Tyr Ala Leu Val Ser Leu Ser Phe Phe Arg Lys Leu
                               265
           260
                                                    270
Arg Leu Ile Arg Gly Glu Thr Leu Glu Ile Gly Asn Tyr Ser Phe Tyr
                           280
                                                285
Ala Leu Asp Asn Gln Asn Leu Arg Gln Leu Trp Asp Trp Ser Lys His
                       295
                                            300
Asn Leu Thr Ile Thr Gln Gly Lys Leu Phe Phe His Tyr Asn Pro Lys
                                       315
                   310
Leu Cys Leu Ser Glu Ile His Lys Met Glu Glu Val Ser Gly Thr Lys
```

```
Gly Arg Gln Glu Arg Asn Asp Ile Ala Leu Lys Thr Asn Gly Asp Gln
                                345
Ala Ser Cys Glu Asn Glu Leu Leu Lys Phe Ser Tyr Ile Arg Thr Ser
        355
                            360
Phe Asp
    370
<210> 105
<211> 383
<212> PRT
<213> Drosophila melanogaster
<400> 105
Arg Gly Gly Val Arg Ile Glu Lys Asn His Lys Leu Cys Tyr Asp Arg
                                     10
Thr Ile Asp Trp Leu Glu Ile Leu Ala Glu Asn Glu Ser Gln Leu Val
            20
Val Leu Thr Glu Asn Gly Lys Glu Lys Glu Cys Ser Leu Ser Lys Cys
Pro Gly Glu Ile Arg Ile Glu Glu Gly His Asp Asn Thr Ala Ile Glu
                        55
Gly Glu Leu Asn Ala Ser Cys Gln Leu His Asn Asn Arg Arg Leu Cys
65
                    70
                                         75
                                                             80
Trp Asn Ser Lys Leu Cys Gln Thr Lys Cys Pro Glu Lys Cys Arg Asn
                85
                                     90
Asn Cys Ile Asp Glu His Thr Cys Cys Ser Gln Asp Cys Leu Gly Gly
            100
                                105
Cys Val Ile Asp Lys Asn Gly Asn Glu Ser Cys Ile Ser Cys Arg Asn
        115
                            120
Val Ser Phe Asn Asn Ile Cys Met Asp Ser Cys Pro Lys Gly Tyr Tyr
                        135
Gln Phe Asp Ser Arg Cys Val Thr Ala Asn Glu Cys Ile Thr Leu Thr
                    150
                                         155
Lys Phe Glu Thr Asn Ser Val Tyr Ser Gly Ile Pro Tyr Asn Gly Gln
                165
                                    170
                                                         175
Cys Ile Thr His Cys Pro Thr Gly Tyr Gln Lys Ser Glu Asn Lys Arg
            180
                                185
Met Cys Glu Pro Cys Pro Gly Gly Lys Cys Asp Lys Glu Cys Ser Ser
        195
                            200
Gly Leu Ile Asp Ser Leu Glu Arg Ala Arg Glu Phe His Gly Cys Thr
                        215
Ile Ile Thr Gly Thr Glu Pro Leu Thr Ile Ser Ile Lys Arg Glu Ser
                    230
                                         235
                                                             240
Gly Ala His Val Met Asp Glu Leu Lys Tyr Gly Leu Ala Ala Val His
                245
                                    250
Lys Ile Gln Ser Ser Leu Met Val His Leu Thr Tyr Gly Leu Lys Ser
                                265
                                                     270
Leu Lys Phe Phe Gln Ser Leu Thr Glu Ile Ser Gly Asp Pro Pro Met
                            280
Asp Ala Asp Lys Tyr Ala Leu Tyr Val Leu Asp Asn Arg Asp Leu Asp
                        295
Glu Leu Trp Gly Pro Asn Gln Thr Val Phe Ile Arg Lys Gly Gly Val
                    310
                                         315
Phe Phe His Phe Asn Pro Lys Leu Cys Val Ser Thr Ile Asn Gln Leu
                                     330
Leu Pro Met Leu Ala Ser Lys Pro Lys Phe Phe Glu Lys Ser Asp Glu
                                 345
Gly Ala Asp Ser Asn Gly Asn Arg Gly Ser Cys Gly Thr Ala Val Leu
```

```
Asn Val Thr Leu Gln Ser Val Gly Ala Asn Ser Ala Ser Leu Asn
    370
                        375
<210> 106
<211> 381
<212> PRT
<213> Caenorhabditis elegans
<400> 106
Asn Gly Gly Val Arg Ile Ile Asp Asn Arg Lys Leu Cys Tyr Thr Lys
Thr Ile Asp Trp Lys His Leu Ile Thr Ser Ser Ile Asn Asp Val Val
            20
Val Asp Asn Ala Ala Glu Tyr Ala Val Thr Glu Thr Gly Leu Met Cys
Pro Arg Gly Ala Cys Glu Glu Asp Lys Gly Glu Ser Lys Cys His Tyr
Leu Glu Glu Lys Asn Gln Glu Gln Gly Val Glu Arg Val Gln Ser Cys
Trp Ser Asn Thr Thr Cys Gln Lys Ser Cys Ala Tyr Asp Arg Leu Leu
                                    90
Pro Thr Lys Glu Ile Gly Pro Gly Cys Asp Ala Asn Gly Asp Arg Cys
                                105
His Asp Gln Cys Val Gly Gly Cys Glu Arg Val Asn Asp Ala Thr Ala
                            120
Cys His Ala Cys Lys Asn Val Tyr His Lys Gly Lys Cys Ile Glu Lys
                        135
Cys Asp Ala His Leu Tyr Leu Leu Gln Arg Arg Cys Val Thr Arg
                    150
                                        155
Glu Gln Cys Leu Gln Leu Asn Pro Val Leu Ser Asn Lys Thr Val Pro
                                    170
Ile Lys Ala Thr Ala Gly Leu Cys Ser Asp Lys Cys Pro Asp Gly Tyr
                                185
Gln Ile Asn Pro Asp Asp His Arg Glu Cys Arg Lys Cys Val Gly Lys
                            200
Cys Glu Ile Val Cys Glu Ile Asn His Val Ile Asp Thr Phe Pro Lys
                        215
Ala Gln Ala Ile Arg Leu Cys Asn Ile Ile Asp Gly Asn Leu Thr Ile
                    230
                                        235
Glu Ile Arg Gly Lys Gln Asp Ser Gly Met Ala Ser Glu Leu Lys Asp
                245
                                    250
Ile Phe Ala Asn Ile His Thr Ile Thr Gly Tyr Leu Leu Val Arg Gln
                                265
Ser Ser Pro Phe Ile Ser Leu Asn Met Phe Arg Asn Leu Arg Arg Ile
Glu Ala Lys Ser Leu Phe Arg Asn Leu Tyr Ala Ile Thr Val Phe Glu
                        295
                                            300
Asn Pro Asn Leu Lys Lys Leu Phe Asp Ser Thr Thr Asp Leu Thr Leu
                    310
                                        315
Asp Arg Gly Thr Val Ser Ile Ala Asn Asn Lys Met Leu Cys Phe Lys
                325
                                    330
Tyr Ile Lys Gln Leu Met Ser Lys Leu Asn Ile Pro Leu Asp Pro Ile
                                345
                                                    350
Asp Gln Ser Glu Gly Thr Asn Gly Glu Lys Ala Ile Cys Glu Asp Met
                            360
Ala Ile Asn Val Ser Ile Thr Ala Val Asn Ala Asp Ser
                        375
```

```
<210> 107
<211> 370
<212> PRT
<213> Homo sapiens
<400> 107
Ala Leu Pro Val Ala Val Leu Leu Ile Val Gly Gly Leu Val Ile Met
1
Leu Tyr Val Phe His Arg Lys Arg Asn Asn Ser Arg Leu Gly Asn Gly
            20
                                25
Val Leu Tyr Ala Ser Val Asn Pro Glu Tyr Phe Ser Ala Ala Asp Val
        35
                            40
Tyr Val Pro Asp Glu Trp Glu Val Ala Arg Glu Lys Ile Thr Met Ser
                        55
Arg Glu Leu Gly Gln Gly Ser Phe Gly Met Val Tyr Glu Gly Val Ala
                    70
Lys Gly Val Val Lys Asp Glu Pro Glu Thr Arg Val Ala Ile Lys Thr
                85
                                    90
Val Asn Glu Ala Ala Ser Met Arg Glu Arg Ile Glu Phe Leu Asn Glu
            100
                                105
Ala Ser Val Met Lys Glu Phe Asn Cys His His Val Val Arg Leu Leu
                            120
Gly Val Val Ser Gln Gly Gln Pro Thr Leu Val Ile Met Glu Leu Met
                        135
                                            140
Thr Arg Gly Asp Leu Lys Ser Tyr Leu Arg Ser Leu Arg Pro Glu Met
                    150
                                        155
Glu Asn Asn Pro Val Leu Ala Pro Pro Ser Leu Ser Lys Met Ile Gln
                165
                                    170
Met Ala Gly Glu Ile Ala Asp Gly Met Ala Tyr Leu Asn Ala Asn Lys
            180
                                185
Phe Val His Arg Asp Leu Ala Ala Arg Asn Cys Met Val Ala Glu Asp
        195
                            200
Phe Thr Val Lys Ile Gly Asp Phe Gly Met Thr Arg Asp Ile Tyr Glu
                        215
Thr Asp Tyr Tyr Arg Lys Gly Gly Lys Gly Leu Leu Pro Val Arg Trp
                    230
                                        235
Met Ser Pro Glu Ser Leu Lys Asp Gly Val Phe Thr Thr Tyr Ser Asp
                245
                                    250
Val Trp Ser Phe Gly Val Val Leu Trp Glu Ile Ala Thr Leu Ala Glu
            260
                                265
Gln Pro Tyr Gln Gly Leu Ser Asn Glu Gln Val Leu Arg Phe Val Met
        275
                            280
Glu Gly Gly Leu Leu Asp Lys Pro Asp Asn Cys Pro Asp Met Leu Phe
    290
                        295
Glu Leu Met Arg Met Cys Trp Gln Tyr Asn Pro Lys Met Arg Pro Ser
                    310
                                        315
Phe Leu Glu Ile Ile Ser Ser Ile Lys Glu Glu Met Glu Pro Gly Phe
                325
                                    330
Arg Glu Val Ser Phe Tyr Tyr Ser Glu Glu Asn Lys Leu Pro Glu Pro
            340
                                345
Glu Glu Leu Asp Leu Glu Pro Glu Asn Met Glu Ser Val Pro Leu Asp
                            360
Pro Ser
    370
<210> 108
<211> 374
```

```
<212> PRT
<213> Homo sapiens
<400> 108
Ile Gly Pro Leu Ile Phe Val Phe Leu Phe Ser Val Val Ile Gly Ser
 1
                                    10
Ile Tyr Leu Phe Leu Arg Lys Arg Gln Pro Asp Gly Pro Leu Gly Pro
            20
                                25
Leu Tyr Ala Ser Ser Asn Pro Glu Tyr Leu Ser Ala Ser Asp Val Phe
Pro Cys Ser Val Tyr Val Pro Asp Glu Trp Glu Val Ser Arg Glu Lys
    50
                        55
Ile Thr Leu Leu Arg Glu Leu Gly Gln Gly Ser Phe Gly Met Val Tyr
                    70
Glu Gly Asn Ala Arg Asp Ile Ile Lys Gly Glu Ala Glu Thr Arg Val
                85
                                    90
Ala Val Lys Thr Val Asn Glu Ser Ala Ser Leu Arg Glu Arg Ile Glu
                                105
Phe Leu Asn Glu Ala Ser Val Met Lys Gly Phe Thr Cys His His Val
                            120
Val Arg Leu Leu Gly Val Val Ser Lys Gly Gln Pro Thr Leu Val Val
                        135
Met Glu Leu Met Ala His Gly Asp Leu Lys Ser Tyr Leu Arg Ser Leu
                                        155
Arg Pro Glu Ala Glu Asn Asn Pro Gly Arg Pro Pro Pro Thr Leu Gln
                                    170
Glu Met Ile Gln Met Ala Ala Glu Ile Ala Asp Gly Met Ala Tyr Leu
Asn Ala Lys Lys Phe Val His Arg Asp Leu Ala Ala Arg Asn Cys Met
        195
                            200
Val Ala His Asp Phe Thr Val Lys Ile Gly Asp Phe Gly Met Thr Arg
                        215
Asp Ile Tyr Glu Thr Asp Tyr Tyr Arg Lys Gly Lys Gly Leu Leu
                    230
                                        235
Pro Val Arg Trp Met Ala Pro Glu Ser Leu Lys Asp Gly Val Phe Thr
                245
                                    250
Thr Ser Ser Asp Met Trp Ser Phe Gly Val Val Leu Trp Glu Ile Thr
            260
                                265
                                                     270
Ser Leu Ala Glu Gln Pro Tyr Gln Gly Leu Ser Asn Glu Gln Val Leu
                            280
Lys Phe Val Met Asp Gly Gly Tyr Leu Asp Gln Pro Asp Asn Cys Pro
                        295
Glu Arg Val Thr Asp Leu Met Arg Met Cys Trp Gln Phe Asn Pro Lys
                    310
                                        315
Met Arg Pro Thr Phe Leu Glu Ile Val Asn Leu Leu Lys Asp Asp Leu
                325
                                    330
His Pro Ser Phe Pro Glu Val Ser Phe Phe His Ser Glu Glu Asn Lys
                                345
Ala Pro Glu Ser Glu Glu Leu Glu Met Glu Phe Glu Asp Met Glu Asn
                            360
Val Pro Leu Asp Arg Ser
    370
<210> 109
<211> 384
```

<212> PRT

<213> Drosophila melanogaster

```
<400> 109
Gly Ile Gly Leu Ala Phe Leu Ile Val Ser Leu Phe Gly Tyr Val Cys
Tyr Leu His Lys Arg Lys Val Pro Ser Asn Asp Leu His Met Asn Thr
            20
Glu Val Asn Pro Phe Tyr Ala Ser Met Gln Tyr Ile Pro Asp Asp Trp
Glu Val Leu Arg Glu Asn Ile Ile Gln Leu Ala Pro Leu Gly Gln Gly
Ser Phe Gly Met Val Tyr Glu Gly Ile Leu Lys Ser Phe Pro Pro Asn
                                        75
Gly Val Asp Arg Glu Cys Ala Ile Lys Thr Val Asn Glu Asn Ala Thr
                                    90
Asp Arg Glu Arg Thr Asn Phe Leu Ser Glu Ala Ser Val Met Lys Glu
            100
                                105
Phe Asp Thr Tyr His Val Val Arg Leu Leu Gly Val Cys Ser Arg Gly
        115
                            120
                                                125
Gln Pro Ala Leu Val Val Met Glu Leu Met Lys Lys Gly Asp Leu Lys
                        135
                                            140
Ser Tyr Leu Arg Ala His Arg Pro Glu Glu Arg Asp Glu Ala Met Met
                    150
                                        155
Thr Tyr Leu Asn Arg Ile Gly Val Thr Gly Asn Val Gln Pro Pro Thr
                                    170
                165
Tyr Gly Arg Ile Tyr Gln Met Ala Ile Glu Ile Ala Asp Gly Met Ala
            180
                                185
                                                    190
Tyr Leu Ala Ala Lys Lys Phe Val His Arg Asp Leu Ala Ala Arg Asn
        195
                            200
Cys Met Val Ala Asp Asp Leu Thr Val Lys Ile Gly Asp Phe Gly Met
                        215
                                            220
Thr Arg Asp Ile Tyr Glu Thr Asp Tyr Tyr Arg Lys Gly Thr Lys Gly
                    230
                                        235
Leu Leu Pro Val Arg Trp Met Pro Pro Glu Ser Leu Arg Asp Gly Val
               245
                                    250
Tyr Ser Ser Ala Ser Asp Val Phe Ser Phe Gly Val Val Leu Trp Glu
                                265
Met Ala Thr Leu Ala Ala Gln Pro Tyr Gln Gly Leu Ser Asn Glu Gln
                            280
                                                285
Val Leu Arg Tyr Val Ile Asp Gly Gly Val Met Glu Arg Pro Glu Asn
                        295
                                            300
Cys Pro Asp Phe Leu His Lys Leu Met Gln Arg Cys Trp His His Arg
                    310
                                        315
Ser Ser Ala Arg Pro Ser Phe Leu Asp Ile Ile Ala Tyr Leu Glu Pro
                                    330
                325
Gln Cys Pro Asn Ser Gln Phe Lys Glu Val Ser Phe Tyr His Ser Glu
            340
                                345
Ala Gly Leu Gln His Arg Glu Lys Glu Arg Lys Glu Arg Asn Gln Leu
                           360
                                                365
Asp Ala Phe Ala Ala Val Pro Leu Asp Gln Asp Leu Gln Asp Arg Glu
                        375
<210> 110
<211> 380
<212> PRT
<213> Caenorhabditis elegans
<400> 110
Gly Met Leu Leu Val Phe Leu Ile Leu Met Ser Ile Ala Gly Cys Ile
```

```
Ile Tyr Tyr Tyr Ile Gln Val Arg Tyr Gly Lys Lys Val Lys Ala Leu
            20
                                 25
Ser Asp Phe Met Gln Leu Asn Pro Glu Tyr Cys Val Asp Asn Lys Tyr
Asn Ala Asp Asp Trp Glu Leu Arg Gln Asp Asp Val Val Leu Gly Gln
                        55
Gln Cys Gly Glu Gly Ser Phe Gly Lys Val Tyr Leu Gly Thr Gly Asn
                                         75
                    70
Asn Val Val Ser Leu Met Gly Asp Arg Phe Gly Pro Cys Ala Ile Lys
                8.5
                                     90
Ile Asn Val Asp Asp Pro Ala Ser Thr Glu Asn Leu Asn Tyr Leu Met
            100
                                 105
                                                     110
Glu Ala Asn Ile Met Lys Asn Phe Lys Thr Asn Phe Ile Val Gln Leu
        115
                            120
                                                 125
Tyr Gly Val Ile Ser Thr Val Gln Pro Ala Met Val Val Met Glu Met
    130
                        135
                                             140
Met Asp Leu Gly Asn Leu Arg Asp Tyr Leu Arg Ser Lys Arg Glu Asp
                    150
                                         155
Glu Val Phe Asn Glu Thr Asp Cys Asn Phe Phe Asp Ile Ile Pro Arg
                165
                                     170
                                                         175
Asp Lys Phe His Glu Trp Ala Ala Gln Ile Cys Asp Gly Met Ala Tyr
            180
                                 185
                                                     190
Leu Glu Ser Leu Lys Phe Cys His Arg Asp Leu Ala Ala Arg Asn Cys
        195
                            200
                                                 205
Met Ile Asn Arg Asp Glu Thr Val Lys Ile Gly Asp Phe Gly Met Ala
                        215
                                             220
Arg Asp Leu Phe Tyr His Asp Tyr Tyr Lys Pro Ser Gly Lys Arg Met
                    230
                                         235
Met Pro Val Arg Trp Met Ser Pro Glu Ser Leu Lys Asp Gly Lys Phe
                245
                                     250
                                                         255
Asp Ser Lys Ser Asp Val Trp Ser Phe Gly Val Val Leu Tyr Glu Met
            260
                                 265
                                                     270
Val Thr Leu Gly Ala Gln Pro Tyr Ile Gly Leu Ser Asn Asp Glu Val
                            280
                                                 285
Leu Asn Tyr Ile Gly Met Ala Arg Lys Val Ile Lys Lys Pro Glu Cys
                        295
                                             300
Cys Glu Asn Tyr Trp Tyr Lys Val Met Lys Met Cys Trp Arg Tyr Ser
                    310
305
                                         315
                                                              320
Pro Arg Asp Arg Pro Thr Phe Leu Gln Leu Val His Leu Leu Ala Ala
                325
                                     330
Glu Ala Ser Pro Glu Phe Arg Asp Leu Ser Phe Val Leu Thr Asp Asn
            340
                                 345
Gln Met Ile Leu Asp Asp Ser Glu Ala Leu Asp Leu Asp Asp Ile Asp
                            360
Asp Thr Asp Met Asn Asp Gln Val Val Glu Val Ala
    370
                        375
```

```
<210> 111
```

<400> 111

Asn Ile Asp Arg Glu Phe Asp Gln Lys Ala Cys Glu Ser Leu Val Lys 1 5 10 15 Lys Leu Lys Asp Lys Lys Asn Asp Leu Gln Asn Leu Ile Asp Val Val 20 25 30 Leu Ser Lys Gly Thr Lys Tyr Thr Gly Cys Ile Thr Ile Pro Arg Thr

<211> 103

<212> PRT

<213> Caenorhabditis elegans

40 Leu Asp Gly Arg Leu Gln Val His Gly Arg Lys Gly Phe Pro His Val Val Tyr Gly Lys Leu Trp Arg Phe Asn Glu Met Thr Lys Asn Glu Thr 70 Arg His Val Asp His Cys Lys His Ala Phe Glu Met Lys Ser Asp Met 90 Val Cys Val Asn Pro Tyr His 100

<210> 112 <211> 104 <212> PRT

<213> Homo sapiens

<400> 112 Gly Gly Glu Ser Glu Thr Phe Ala Lys Arg Ala Ile Glu Ser Leu Val Lys Lys Leu Lys Glu Lys Lys Asp Glu Leu Asp Ser Leu Ile Thr Ala 25 Ile Thr Thr Asn Gly Ala His Pro Ser Lys Cys Val Thr Ile Gln Arg 40 Thr Leu Asp Gly Arg Leu Gln Val Ala Gly Arg Lys Gly Phe Pro His 55 Val Ile Tyr Ala Arg Leu Trp Arg Trp Pro Asp Leu His Lys Asn Glu 75 Leu Lys His Val Lys Tyr Cys Gln Tyr Ala Phe Asp Leu Lys Cys Asp 85 Ser Val Cys Val Asn Pro Tyr His 100

<210> 113 <211> 205 <212> PRT <213> Caenorhabditis elegans

<400> 113 Ile Val Tyr Tyr Glu Lys Asn Leu Gln Ile Gly Glu Lys Lys Cys Ser Arg Gly Asn Phe His Val Asp Gly Gly Phe Ile Cys Ser Glu Asn Arg 25 Tyr Ser Leu Gly Leu Glu Pro Asn Pro Ile Arg Glu Pro Val Ala Phe 40 Lys Val Arg Lys Ala Ile Val Asp Gly Ile Arg Phe Ser Tyr Lys Lys 55 Asp Gly Ser Val Trp Leu Gln Asn Arg Met Lys Tyr Pro Val Phe Val 70 75 Thr Ser Gly Tyr Leu Asp Glu Gln Ser Gly Gly Leu Lys Lys Asp Lys 85 90 Val His Lys Val Tyr Gly Cys Ala Ser Ile Lys Thr Phe Gly Phe Asn 100 105 110 Val Ser Lys Gln Ile Ile Arg Asp Ala Leu Leu Ser Lys Gln Met Ala 115 120 125 Thr Met Tyr Leu Gln Gly Lys Leu Thr Pro Met Asn Tyr Ile Tyr Glu 135 140 Lys Lys Thr Gln Glu Glu Leu Arg Arg Glu Ala Thr Arg Thr Thr Asp 150 155

```
Ser Leu Ala Lys Tyr Cys Cys Val Arg Val Ser Phe Cys Lys Gly Phe
                165
                                    170
Gly Glu Ala Tyr Pro Glu Arg Pro Ser Ile His Asp Cys Pro Val Trp
            180
                                185
Ile Glu Leu Lys Ile Asn Ile Ala Tyr Asp Phe Met Asp
<210> 114
<211> 212
<212> PRT
<213> Homo sapiens
<400> 114
Ile Ala Tyr Phe Glu Met Asp Val Gln Val Gly Glu Thr Phe Lys Val
                                                       15
                                    10
Pro Ser Ser Cys Pro Ile Val Thr Val Asp Gly Tyr Val Asp Pro Ser
            20
                                25
                                                    30
Gly Gly Asp Arg Phe Cys Leu Gly Gln Leu Ser Asn Val His Arg Thr
                            40
Glu Ala Ile Glu Arg Ala Arg Leu His Ile Gly Lys Gly Val Gln Leu
                        55
                                            60
Glu Cys Lys Gly Glu Gly Asp Val Trp Val Arg Cys Leu Ser Asp His
                   70
                                        75
Ala Val Phe Val Gln Ser Tyr Tyr Leu Asp Arg Glu Ala Gly Arg Ala
                                    90
               85
Pro Gly Asp Ala Val His Lys Ile Tyr Pro Ser Ala Tyr Ile Lys Val
           100
                                105
Phe Asp Leu Arg Gln Cys His Arg Gln Met Gln Gln Ala Ala Thr
       115
                           120
                                                125
Ala Gln Ala Ala Ala Ala Gln Ala Ala Val Ala Gly Asn Ile
                       135
                                            140
Pro Gly Pro Gly Ser Val Gly Gly Ile Ala Pro Ala Ile Ser Leu Ser
                   150
                                        155
Ala Ala Ala Gly Ile Gly Val Asp Asp Leu Arg Arg Leu Cys Ile Leu
                                    170
               165
Arg Met Ser Phe Val Lys Gly Trp Gly Pro Asp Tyr Pro Arg Gln Ser
                               185
                                                    190
Ile Lys Glu Thr Pro Cys Trp Ile Glu Ile His Leu His Arg Ala Leu
       195
                           200
Gln Leu Leu Asp
   210
<210> 115
<211> 50
<212> PRT
<213> Caenorhabditis elegans
<220>
<221> VARIANT
<222> (1)...(50)
<223> Xaa = Any Amino Acid
<400> 115
Leu Cys Gly Xaa Xaa Leu Val Glu Ala Leu Xaa Xaa Val Cys Gly Xaa
                                    10
Arg Gly Phe Phe Tyr Thr Pro Lys Thr Arg Arg Lys Arg Gly Ile Val
```

```
Glu Gln Cys Cys Xaa Xaa Xaa Cys Xaa Xaa Xaa Gln Leu Glu Xaa Tyr
                            40
Cys Asn
    50
<210> 116
<211> 39
<212> PRT
<213> Caenorhabditis elegans
<400> 116
Leu Cys Gly Arg His Leu Ala Asp Ala Leu Tyr Phe Val Cys Gly Asn
Arg Gly Phe Gly Ile Val Glu Glu Cys Cys His Asn Pro Cys Thr Leu
                                25
Tyr Gln Leu Glu Asn Tyr Cys
        35
<210> 117
<211> 112
<212> PRT
<213> Caenorhabditis elegans
<400> 117
Met Asn Ser Val Phe Thr Ile Ile Phe Val Leu Cys Ala Leu Gln Val
Ala Ala Ser Phe Arg Gln Ser Phe Gly Pro Ser Met Ser Glu Glu Ser
                                25
Ala Ser Met Gln Leu Leu Arg Glu Leu Gln His Asn Met Met Glu Ser
Ala His Arg Pro Met Pro Arg Ala Arg Arg Val Pro Ala Pro Gly Glu
Thr Arg Ala Cys Gly Arg Lys Leu Ile Ser Leu Val Met Ala Val Cys
                    70
Gly Asp Leu Cys Asn Pro Gln Glu Gly Lys Asp Ile Ala Thr Glu Cys
                                    90
Cys Gly Asn Gln Cys Ser Asp Asp Tyr Ile Arg Ser Ala Cys Cys Pro
                                105
<210> 118
<211> 106
<212> PRT
<213> Caenorhabditis elegans
<400> 118
Met Phe Ser Phe Phe Thr Tyr Phe Leu Leu Ser Ala Leu Leu Ser
                                    10
Ala Ser Cys Arg Gln Pro Ser Met Asp Thr Ser Lys Ala Asp Arg Ile
                                25
Leu Arg Glu Ile Glu Met Glu Thr Glu Leu Glu Asn Gln Leu Ser Arg
Ala Arg Arg Val Pro Ala Gly Glu Val Arg Ala Cys Gly Arg Arg Leu
                        55
Leu Leu Phe Val Trp Ser Thr Cys Gly Glu Pro Cys Thr Pro Gln Glu
                    70
                                        75
Asp Met Asp Ile Ala Thr Val Cys Cys Thr Thr Gln Cys Thr Pro Ser
```

```
95
                85
Tyr Ile Lys Gln Ala Cys Cys Pro Glu Lys
            100
                                105
<210> 119
<211> 105
<212> PRT
<213> Caenorhabditis elegans
<400> 119
Met Pro Pro Ile Ile Leu Val Phe Phe Leu Val Leu Ile Pro Ala Ser
Gln Gln Tyr Pro Phe Ser Leu Glu Ser Leu Asn Asp Gln Ile Ile Asn
Glu Glu Val Ile Glu Tyr Met Leu Glu Asn Ser Ile Arg Ser Ser Arg
Thr Arg Arg Val Pro Asp Glu Lys Lys Ile Tyr Arg Cys Gly Arg Arg
Ile His Ser Tyr Val Phe Ala Val Cys Gly Lys Ala Cys Glu Ser Asn
                    70
Thr Glu Val Asn Ile Ala Ser Lys Cys Cys Arg Glu Glu Cys Thr Asp
                                    90
Asp Phe Ile Arg Lys Gln Cys Cys Pro
            100
<210> 120
<211> 118
<212> PRT
<213> Caenorhabditis elegans
<400> 120
Met Ile Val Thr Leu Ile Val Phe Leu Val Ile Gly Leu Gln Met Ala
His Leu Ser Gln Val Ser Gly Asn Asn Glu Asn Gly Phe Leu Asn Pro
            20
                                25
Phe Asp Leu Ser Gln Trp Ser Glu Glu Ile Leu His Arg Gln Tyr His
                            40
His His His His His His Gly Asn Arg Ala Arg Thr Leu Glu
Thr Glu Lys Ile Tyr Arg Cys Gly Arg Lys Leu Tyr Thr Asp Val Leu
                    70
Ser Ala Cys Asn Gly Pro Cys Glu Pro Gly Thr Glu Gln Asp Leu Ser
                                    90
Lys Leu Cys Cys Gly Asn Gln Cys Thr Phe Val Glu Ile Arg Lys Ala
                                105
Cys Cys Ala Asp Lys Leu
        115
<210> 121
<211> 106
<212> PRT
<213> Caenorhabditis elegans
<400> 121
Met Asn Ala Ile Ile Phe Cys Leu Leu Phe Thr Thr Val Thr Ala Thr
```

```
Tyr Glu Val Phe Gly Lys Gly Ile Glu His Arg Asn Glu His Leu Ile
Ile Asn Gln Leu Asp Ile Ile Pro Val Glu Ser Thr Pro Thr Pro Asn
Arg Ala Ser Arg Val Gln Lys Arg Leu Cys Gly Arg Arg Leu Ile Leu
Phe Met Leu Ala Thr Cys Gly Glu Cys Asp Thr Asp Ser Ser Glu Asp
                                        75
                                                            80
Leu Ser His Ile Cys Cys Ile Lys Gln Cys Asp Val Gln Asp Ile Ile
                85
                                    90
Arg Val Cys Cys Pro Asn Ser Phe Arg Lys
           100
<210> 122
```

<211> 107 <212> PRT <213> Caenorhabditis elegans

<400> 122

Met Lys Leu Ser Val Val Leu Ala Leu Phe Ile Ile Phe Gln Leu Gly 10 Ala Ala Ser Leu Met Arg Asn Trp Met Phe Asp Phe Glu Lys Glu Leu 25 30 Glu His Asp Tyr Asp Asp Ser Glu Ile Gly Phe His Asn Ile His Ser 35 40 Leu Met Ala Arg Ser Arg Arg Gly Asp Lys Val Lys Ile Cys Gly Thr 55 60 Lys Val Leu Lys Met Val Met Val Met Cys Gly Glu Cys Ser Ser 65 70 75 Thr Asn Glu Asn Ile Ala Thr Glu Cys Cys Glu Lys Met Cys Thr Met 85 90 Glu Asp Ile Thr Thr Lys Cys Cys Pro Ser Arg

<210> 123 <211> 73 <212> PRT <213> Caenorhabditis elegans

<400> 123 Met Lys Leu His Ile Phe Ile Ile Phe Leu Leu Phe Gln Ser Cys 10 Ser Asn Lys Met Cys Gln Tyr Ser Lys Lys Tyr Lys Ile Cys Gly 20 25 Val Arg Ala Leu Lys His Met Lys Val Tyr Cys Thr Arg Gly Met Thr 40 Arg Asp Tyr Gly Lys Leu Leu Val Thr Cys Cys Ser Lys Gly Cys Asn 55 Ala Ile Asp Ile Gln Arg Ile Cys Leu

<210> 124 <211> 109 <212> PRT <213> Caenorhabditis elegans

```
<400> 124
Met Tyr Trp Phe Arg Gln Val Tyr Arg Pro Ser Phe Phe Phe Gly Phe
                                    10
Leu Ala Ile Leu Leu Ser Ser Pro Thr Pro Ser Asp Ala Ser Ile
                                25
Arg Leu Cys Gly Ser Arg Leu Thr Thr Leu Leu Ala Val Cys Arg
                            40
Asn Gln Leu Cys Thr Gly Leu Thr Ala Phe Lys Arg Ser Ala Asp Gln
                        5.5
Ser Tyr Ala Pro Thr Thr Arg Asp Leu Phe His Ile His His Gln Gln
                    70
                                        75
Lys Arg Gly Gly Ile Ala Thr Glu Cys Cys Glu Lys Arg Cys Ser Phe
                                    90
                85
Ala Tyr Leu Lys Thr Phe Cys Cys Asn Gln Asp Asp Asn
            100
                                105
<210> 125
<211> 110
<212> PRT
<213> Homo sapiens
<400> 125
Met Ala Leu Trp Met Arg Leu Leu Pro Leu Leu Ala Leu Leu Ala Leu
                                    10
Trp Gly Pro Asp Pro Ala Ala Ala Phe Val Asn Gln His Leu Cys Gly
                                25
Ser His Leu Val Glu Ala Leu Tyr Leu Val Cys Gly Glu Arg Gly Phe
                            40
                                                45
Phe Tyr Thr Pro Lys Thr Arg Arg Glu Ala Glu Asp Leu Gln Val Gly
                        55
                                            60
Gln Val Glu Leu Gly Gly Gly Pro Gly Ala Gly Ser Leu Gln Pro Leu
                   70
                                        75
Ala Leu Glu Gly Ser Leu Gln Lys Arg Gly Ile Val Glu Gln Cys Cys
                                   90
Thr Ser Ile Cys Ser Leu Tyr Gln Leu Glu Asn Tyr Cys Asn
                                105
<210> 126
<211> 46
<212> PRT
<213> Caenorhabditis elegans
<220>
<221> VARIANT
<222> (1)...(46)
<223> Xaa = Any Amino Acid
<400> 126
Ala Cys Gly Arg Arg Leu Leu Phe Val Trp Ser Thr Cys Gly Glu
                                   10
Pro Cys Thr Xaa Xaa Xaa Gln Glu Asp Met Asp Ile Ala Thr Val Cys
                                25
Cys Thr Thr Gln Cys Thr Pro Ser Tyr Ile Lys Gln Ala Cys
```

```
<211> 46
<212> PRT
<213> Caenorhabditis elegans
<220>
<221> VARIANT
<222> (1)...(46)
<223> Xaa = Any Amino Acid
<400> 127
Ala Cys Gly Arg Lys Leu Ile Ser Leu Val Met Ala Val Cys Gly Asp
1
                                                         15
Leu Cys Asn Xaa Xaa Xaa Gln Glu Gly Lys Asp Ile Ala Thr Glu Cys
            20
                                 25
Cys Gly Asn Gln Cys Ser Asp Asp Tyr Ile Arg Ser Ala Cys
                            40
<210> 128
<211> 46
<212> PRT
<213> Caenorhabditis elegans
<220>
<221> VARIANT
<222> (1)...(46)
<223> Xaa = Any Amino Acid
<400> 128
Arg Cys Gly Arg Arg Ile His Ser Tyr Val Phe Ala Val Cys Gly Lys
                                     10
Ala Cys Glu Xaa Xaa Ser Thr Glu Val Asn Ile Ala Ser Lys Cys
            20
                                 25
Cys Arg Glu Glu Cys Thr Asp Asp Phe Ile Arg Lys Gln Cys
        35
                            40
<210> 129
<211> 46
<212> PRT
<213> Caenorhabditis elegans
<220>
<221> VARIANT
<222> (1) ... (46)
<223> Xaa = Any Amino Acid
<400> 129
Arg Cys Gly Arg Lys Leu Tyr Thr Asp Val Leu Ser Ala Cys Asn Gly
1
                                     10
Pro Cys Glu Xaa Xaa Xaa Gly Thr Glu Gln Asp Leu Ser Lys Leu Cys
                                 25
                                                     30
Cys Gly Asn Gln Cys Thr Phe Asx Glu Ile Arg Lys Ala Cys
<210> 130
<211> 46
<212> PRT
```

```
<213> Caenorhabditis elegans
<220>
<221> VARIANT
<222> (1)...(46)
<223> Xaa = Any Amino Acid
<400> 130
Ile Cys Gly Thr Lys Asx Leu Lys Met Val Met Val Met Cys Gly Gly
1
                                     10
                                                         15
Glu Cys Ser Xaa Xaa Xaa Ser Thr Asn Glu Asn Ile Ala Thr Glu Cys
            20
                                 25
Cys Glu Lys Met Cys Thr Met Glu Asp Ile Thr Thr Lys Cys
        35
<210> 131
<211> 46
<212> PRT
<213> Caenorhabditis elegans
<220>
<221> VARIANT
<222> '(1) ... (46)
<223> Xaa = Any Amino Acid
<400> 131
Leu Cys Gly Arg Arg Leu Ile Leu Phe Met Leu Ala Thr Cys Gly Glu
                                     10
Cys Asp Thr Xaa Xaa Xaa Asp Ser Ser Glu Asp Leu Ser His Ile Cys
            20
                                 25
Cys Ile Lys Gln Cys Asp Val Gln Asp Ile Ile Arg Val Cys
        35
                             40
<210> 132
<211> 46
<212> PRT
<213> Caenorhabditis elegans
<220>
<221> VARIANT
<222> (1)...(46)
<223> Xaa = Any Amino Acid
<400> 132
Leu Cys Gly Ser His Leu Val Glu Ala Leu Tyr Leu Val Cys Gly Glu
1
                                                         15
Arg Gly Phe Xaa Xaa Xaa Leu Gln Lys Arg Gly Ile Val Glu Gln Cys
                                 25
            20
Cys Thr Ser Ile Cys Ser Leu Tyr Gln Leu Glu Asn Tyr Cys
        35
                             40
<210> 133
<211> 46
<212> PRT
<213> Rabbit
```

```
<220>
<221> VARIANT
<222> (1)...(46)
<223> Xaa = Any Amino Acid
<400> 133
Leu Cys Gly Ser His Leu Val Glu Ala Leu Tyr Leu Val Cys Gly Glu
1
                 5
Arg Gly Phe Xaa Xaa Xaa Thr Pro Lys Ser Gly Ile Val Glu Gln Cys
                                25
            20
Cys Thr Ser Ile Cys Ser Leu Tyr Gln Leu Glu Asn Tyr Cys
                            40
                                                 45
<210> 134
<211> 46
<212> PRT
<213> Xenopus laevis
<220>
<221> VARIANT
<222> (1) ... (46)
<223> Xaa = Any Amino Acid
<400> 134
Leu Cys Gly Ser His Leu Val Glu Ala Leu Tyr Leu Val Cys Gly Asp
1
Arg Gly Phe Xaa Xaa Lys Met Lys Arg Gly Ile Val Glu Gln Cys
                                25
            20
Cys His Ser Thr Cys Ser Leu Phe Gln Leu Glu Ser Tyr Cys
        35
                            40
<210> 135
<211> 46
<212> PRT
<213> Xenopus laevis
<220>
<221> VARIANT
<222> (1)...(46)
<223> Xaa = Any Amino Acid
<400> 135
Leu Cys Gly Ser His Leu Val Glu Ala Leu Tyr Leu Val Cys Gly Asp
                 5
Arg Gly Phe Xaa Xaa Xaa Lys Met Lys Arg Gly Ile Val Glu Gln Cys
            20
                                25
Cys His Ser Thr Cys Ser Leu Phe Gln Leu Glu Asn Tyr Cys
                                                 45
<210> 136
<211> 46
<212> PRT
<213> Alligator
<220>
<221> VARIANT
```

```
<222> (1)...(46)
<223> Xaa = Any Amino Acid
<400> 136
Leu Cys Gly Ser His Leu Val Asp Ala Leu Tyr Leu Val Cys Gly Glu
                 5
                                                          15
Arg Gly Phe Xaa Xaa Xaa Ser Pro Lys Gly Gly Ile Val Glu Gln Cys
            20
                                 25
Cys His Asn Thr Cys Ser Leu Tyr Gln Leu Glu Asn Tyr Cys
                             40
                                                 45
<210> 137
<211> 46
<212> PRT
<213> Elephant fish
<220>
<221> VARIANT
<222> (1)...(46)
<223> Xaa = Any Amino Acid
<400> 137
Leu Cys Gly Ser His Leu Val Asp Ala Leu Tyr Phe Val Cys Gly Glu
Arg Gly Phe Xaa Xaa Xaa Pro Lys Gln Ile Gly Ile Val Glu Gln Cys
            20
                                 25
                                                      30
Cys His Asn Thr Cys Ser Leu Val Asn Leu Glu Gly Tyr Cys
                             40
                                                 45
<210> 138
<211> 46
<212> PRT
<213> Bos taurus
<220>
<221> VARIANT
<222> (1)...(46)
<223> Xaa = Any Amino Acid
<400> 138
Leu Cys Gly Ala Glu Leu Val Asp Ala Leu Gln Phe Val Cys Gly Asp
                 5
                                     10
                                                          15
Arg Gly Phe Xaa Xaa Xaa Ala Pro Gln Thr Gly Ile Val Asp Glu Cys
                                 25
Cys Phe Arg Ser Cys Asp Leu Arg Arg Leu Glu Met Tyr Cys
                                                 45
                             40
<210> 139
<211> 46
<212> PRT
<213> Canis
<220>
<221> VARIANT
<222> (1)...(46)
```

<223> Xaa = Any Amino Acid

```
<400> 139
Leu Cys Gly Ala Glu Leu Val Asp Ala Leu Gln Phe Val Cys Gly Asp
Arg Gly Phe Xaa Xaa Xaa Ala Pro Gln Thr Gly Ile Val Asp Glu Cys
Cys Phe Arg Ser Cys Asp Leu Arg Arg Leu Glu Met Tyr Cys
<210> 140
<211> 46
<212> PRT
<213> Horse
<220>
<221> VARIANT
<222> (1)...(46)
<223> Xaa = Any Amino Acid
<400> 140
Leu Cys Gly Glu Leu Val Asp Thr Leu Gln Phe Val Cys Gly Asp
1
                 5
                                                         15
                                    10
Arg Gly Phe Xaa Xaa Xaa Arg Arg Ser Arg Gly Ile Val Glu Glu Cys
                                25
Cys Phe Arg Ser Cys Asp Leu Ala Leu Leu Glu Thr Tyr Cys
        35
<210> 141
<211> 46
<212> PRT
<213> Homo sapiens
<220>
<221> VARIANT
<222> (1) ... (46)
<223> Xaa = Any Amino Acid
<400> 141
Leu Cys Gly Glu Leu Val Asp Thr Leu Gln Phe Val Cys Gly Asp
                                     10
Arg Gly Phe Xaa Xaa Xaa Arg Arg Ser Arg Gly Ile Val Glu Cys
                                25
Cys Phe Arg Ser Cys Asp Leu Ala Leu Leu Glu Thr Tyr Cys
                            40
<210> 142
<211> 46
<212> PRT
<213> Amphioxus
<220>
<221> VARIANT
<222> (1) ... (46)
<223> Xaa = Any Amino Acid
<400> 142
Leu Cys Gly Ser Thr Leu Ala Asp Val Leu Ser Phe Val Cys Gly Asn
```

```
10
Arg Gly Tyr Xaa Xaa Xaa Arg Arg Arg Gly Leu Val Glu Glu Cys
            20
                                25
Cys Tyr Asn Val Cys Asp Tyr Ser Gln Leu Glu Ser Tyr Cys
                             40
<210> 143
<211> 46
<212> PRT
<213> Locust
<220>
<221> VARIANT
<222> (1)...(46)
<223> Xaa = Any Amino Acid
<400> 143
Tyr Cys Gly Glu Lys Leu Ser Asn Ala Leu Lys Leu Val Cys Arg Gly
1
                                                         15
Asn Tyr Asn Xaa Xaa Xaa Arg Arg Thr Arg Gly Val Phe Asp Glu Cys
                                25
            20
Cys Arg Lys Ser Cys Ser Ile Ser Glu Leu Gln Thr Tyr Cys
        35
                                                 45
<210> 144
<211> 46
<212> PRT
<213> Bommo
<220>
<221> VARIANT
<222> (1)...(46)
<223> Xaa = Any Amino Acid
<400> 144
Tyr Cys Gly Arg His Leu Ala Arg Thr Leu Ala Asp Leu Cys Trp Glu
                                    10
Ala Gly Val Xaa Xaa Xaa Arg Gly Lys Arg Gly Ile Val Asp Glu Cys
                                25
            20
Cys Leu Arg Pro Cys Ser Val Asp Val Leu Leu Ser Tyr Cys
        35
                            40
<210> 145
<211> 46
<212> PRT
<213> Bommo
<220>
<221> VARIANT
<222> (1)...(46)
<223> Xaa = Any Amino Acid
<400> 145
Tyr Cys Gly Arg His Leu Ala Asp Thr Leu Ala Asp Leu Cys Phe Gly
1
                                    10
Val Glu Lys Xaa Xaa Xaa Arg Gly Lys Arg Gly Val Val Asp Glu Cys
```

```
25
Cys Phe Arg Pro Cys Thr Leu Asp Val Leu Leu Ser Tyr Cys
        35
                             40
<210> 146
<211> 46
<212> PRT
<213> Horn worm
<220>
<221> VARIANT
<222> (1)...(46)
<223> Xaa = Any Amino Acid
<400> 146
Ile Cys Gly Arg His Leu Ala Arg Thr Leu Ala Asp Leu Cys Pro Asn
1
                                     10
Val Glu Tyr Xaa Xaa Xaa Gly Lys Arg Ala Gly Val Ala Asp Asp Cys
            20
                                 25
Cys Asx Asn Ser Cys Thr Met Asp Val Leu Leu Ser Tyr Cys
                             40
                                                 45
<210> 147
<211> 46
<212> PRT
<213> Bombyx mori
<220>
<221> VARIANT
<222> (1)...(46)
<223> Xaa = Any Amino Acid
<400> 147
Tyr Cys Gly Arg Arg Leu Ala Thr Met Leu Ser Phe Val Cys Asp Asn
                                     10
Gln Tyr Gln Xaa Xaa Xaa Gly Lys Arg Gln Gly Ile Ala Glu Glu Cys
            20
                                 25
Cys Asn Lys Pro Cys Thr Glu Asn Glu Leu Leu Gly Tyr Cys
        35
                             40
<210> 148
<211> 46
<212> PRT
<213> Bombyx mori
<220>
<221> VARIANT
<222> (1)...(46)
<223> Xaa = Any Amino Acid
<400> 148
Tyr Cys Gly Arg Arg Leu Ala Thr Met Leu Leu Tyr Val Cys Asp Asn
1
                 5
                                     10
                                                         15
Gln Tyr Gln Xaa Xaa Xaa Gly Lys Arg Gln Gly Ile Val Glu Glu Cys
            20
                                 25
Cys Asn Lys Pro Cys Thr Glu Asn Glu Leu Leu Gly Tyr Cys
```

```
<210> 149
<211> 46
<212> PRT
<213> Bombys mori
<220>
<221> VARIANT
<222> (1)...(46)
<223> Xaa = Any Amino Acid
<400> 149
Tyr Cys Gly Arg Arg Leu Ala Ile Met Leu Ser Tyr Leu Cys Asp Asn
1
Gln Tyr Leu Xaa Xaa Xaa Gly Lys Arg Gln Gly Ile Ala Glu Glu Cys
            20
                                 25
Cys Asn Lys Pro Cys Thr Glu Asp Glu Leu Leu Gly Tyr Cys
        35
                             40
<210> 150
<211> 46
<212> PRT
<213> Caenorhabditis elegans
<220>
<221> VARIANT
<222> (1) ... (46)
<223> Xaa = Any Amino Acid
<400> 150
Leu Cys Gly Ser Arg Leu Thr Thr Thr Leu Leu Ala Val Cys Arg Asn
Gln Leu Cys Xaa Xaa Xaa Gln Lys Arg Gly Gly Ile Ala Thr Glu Cys
            20
                                 25
Cys Glu Lys Arg Cys Ser Phe Ala Tyr Leu Lys Thr Phe Cys
        35
                             40
<210> 151
<211> 46
<212> PRT
<213> Moi 3
<220>
<221> VARIANT
<222> (1) ... (46)
<223> Xaa = Any Amino Acid
<400> 151
Leu Cys Gly Ser Thr Leu Ala Asn Met Val Gln Trp Leu Cys Ser Thr
                 5
                                     10
Tyr Thr Thr Xaa Xaa Xaa Glu Ser Arg Pro Ser Ile Val Cys Glu Cys
                                 25
                                                     30
Cys Phe Asn Gln Cys Thr Val Gln Glu Leu Leu Ala Tyr Cys
```

```
<210> 152
<211> 46
<212> PRT
<213> Homo sapiens
<220>
<221> VARIANT
<222> (1)...(46)
<223> Xaa = Any Amino Acid
<400> 152
Leu Cys Gly Arg Glu Leu Val Arg Ala Gln Ile Ala Ile Cys Gly Met
1
                                     10
Ser Thr Trp Xaa Xaa Xaa Arg Pro Tyr Val Ala Leu Phe Glu Lys Cys
            20
                                 25
Cys Leu Ile Gly Cys Thr Lys Arg Ser Leu Ala Lys Tyr Cys
                            40
<210> 153
<211> 46
<212> PRT
<213> Homo sapiens
<220>
<221> VARIANT
<222> (1)...(46)
<223> Xaa = Any Amino Acid
<400> 153
Leu Cys Gly His His Phe Val Arg Ala Leu Val Arg Val Cys Gly Gly
                                                         15
                                     10
Pro Arg Trp Xaa Xaa Xaa Ala Ala Ala Thr Asn Pro Ala Arg Tyr Cys
            20
                                 25
Cys Leu Ser Gly Cys Thr Gln Gln Asp Leu Leu Thr Leu Cys
        35
<210> 154
<211> 541
<212> PRT
<213> Caenorhabditis elegans
<400> 154
Met Ser Met Thr Ser Leu Ser Thr Lys Ser Arg Arg Gln Glu Asp Val
1
Val Ile Glu Gly Trp Leu His Lys Lys Gly Glu His Ile Arg Asn Trp
                                25
Arg Pro Arg Tyr Phe Met Ile Phe Asn Asp Gly Ala Leu Leu Gly Phe
        35
Arg Ala Lys Pro Lys Glu Gly Gln Pro Phe Pro Glu Pro Leu Asn Asp
                        55
Phe Met Ile Lys Asp Ala Ala Thr Met Leu Phe Glu Lys Pro Arg Pro
                    70
Asn Met Phe Met Val Arg Cys Leu Gln Trp Thr Thr Val Ile Glu Arg
                85
                                     90
Thr Phe Tyr Ala Glu Ser Ala Glu Val Arg Gln Arg Trp Ile His Ala
            100
                                105
Ile Glu Ser Ile Ser Lys Lys Tyr Lys Gly Thr Asn Ala Asn Pro Gln
```

```
120
       115
Glu Glu Leu Met Glu Thr Asn Gln Gln Pro Lys Ile Asp Glu Asp Ser
                      135
                                           140
Glu Phe Ala Gly Ala Ala His Ala Ile Met Gly Gln Pro Ser Ser Gly
                                      155
                   150
His Gly Asp Asn Cys Ser Ile Asp Phe Arg Ala Ser Met Ile Ser Ile
                                  170
              165
Ala Asp Thr Ser Glu Ala Ala Lys Arg Asp Lys Ile Thr Met Glu Asp
                              185
Phe Asp Phe Leu Lys Val Leu Gly Lys Gly Thr Phe Gly Lys Val Ile
                           200
                                               205
Leu Cys Lys Glu Lys Arg Thr Gln Lys Leu Tyr Ala Ile Lys Ile Leu
                                           220
                      215
Lys Lys Asp Val Ile Ile Ala Arg Glu Val Ala His Thr Leu Thr
                  230
                                      235
Glu Asn Arg Val Leu Gln Arg Cys Lys His Pro Phe Leu Thr Glu Leu
              245
                                   250
Lys Tyr Ser Phe Gln Glu Gln His Tyr Leu Cys Phe Val Met Gln Phe
                              265
Ala Asn Gly Gly Glu Leu Phe Thr His Val Arg Lys Cys Gly Thr Phe
                           280
                                               285
Ser Glu Pro Arg Ala Arg Phe Tyr Gly Ala Glu Ile Val Leu Ala Leu
                      295
                                          300
Gly Tyr Leu His Arg Cys Asp Ile Val Tyr Arg Asp Met Lys Leu Glu
                  310
                                       315
Asn Leu Leu Leu Asp Lys Asp Gly His Ile Lys Ile Ala Asp Phe Gly
              325
                                   330
Leu Cys Lys Glu Glu Ile Ser Phe Gly Asp Lys Thr Ser Thr Phe Cys
           340
                              345
Gly Thr Pro Glu Tyr Leu Ala Pro Glu Val Leu Asp Asp His Asp Tyr
                          360
                                               365
Gly Arg Cys Val Asp Trp Trp Gly Val Gly Val Val Met Tyr Glu Met
                      375
                                          380
Met Cys Gly Arg Leu Pro Phe Tyr Ser Lys Asp His Asn Lys Leu Phe
                  390
                                      395
Glu Leu Ile Met Ala Gly Asp Leu Arg Phe Pro Ser Lys Leu Ser Gln
               405
                                   410
Glu Ala Arg Thr Leu Leu Thr Gly Leu Leu Val Lys Asp Pro Thr Gln
           420
                              425
                                                  430
Arg Leu Gly Gly Gly Pro Glu Asp Ala Leu Glu Ile Cys Arg Ala Asp
                          440
                                              445
Phe Phe Arg Thr Val Asp Trp Glu Ala Thr Tyr Arg Lys Glu Ile Glu
                      455
                                           460
Pro Pro Tyr Lys Pro Asn Val Gln Ser Glu Thr Asp Thr Ser Tyr Phe
                   470
                                       475
Asp Asn Glu Phe Thr Ser Gln Pro Val Gln Leu Thr Pro Pro Ser Arg
               485
                                   490
Ser Gly Ala Leu Ala Thr Val Asp Glu Glu Glu Met Gln Ser Asn
                               505
Phe Thr Gln Phe Ser Phe His Asn Val Met Gly Ser Ile Asn Arg Ile
                          520
His Glu Ala Ser Glu Asp Asn Glu Asp Tyr Asp Met Gly
                       535
```

<210> 155

<211> 546

<212> PRT

<213> Caenorhabditis elegans

<400> 155 Met Ser Met Thr Ser Leu Ser Thr Lys Ser Arg Arg Gln Glu Asp Val Val Ile Glu Gly Trp Leu His Lys Lys Gly Glu His Ile Arg Asn Trp 25 Arg Pro Arg Tyr Phe Met Ile Phe Asn Asp Gly Ala Leu Leu Gly Phe Arg Ala Lys Pro Lys Glu Gly Gln Pro Phe Pro Glu Pro Leu Asn Asp 55 Phe Met Ile Lys Asp Ala Ala Thr Met Leu Phe Glu Lys Pro Arg Pro 70 7.5 Asn Met Phe Met Val Arg Cys Leu Gln Trp Thr Thr Val Ile Glu Arg 8.5 90 Thr Phe Tyr Ala Glu Ser Ala Glu Val Arg Gln Arg Trp Ile His Ala 100 105 Ile Glu Ser Ile Ser Lys Lys Tyr Lys Gly Thr Asn Ala Asn Pro Gln 120 125 Glu Glu Leu Met Glu Thr Asn Gln Gln Pro Lys Ile Asp Glu Asp Ser 135 140 Glu Phe Ala Gly Ala Ala His Ala Ile Met Gly Gln Pro Ser Ser Gly 150 155 His Gly Asp Asn Cys Ser Ile Asp Phe Arg Ala Ser Met Ile Ser Ile 170 165 Ala Asp Thr Ser Glu Ala Ala Lys Arg Asp Lys Ile Thr Met Glu Asp 180 185 Phe Asp Phe Leu Lys Val Leu Gly Lys Gly Thr Phe Gly Lys Val Ile 200 205 Leu Cys Lys Glu Lys Arg Thr Gln Lys Leu Tyr Ala Ile Lys Ile Leu 215 220 Lys Lys Asp Val Ile Ile Ala Arg Glu Val Ala His Thr Leu Thr 230 235 Glu Asn Arg Val Leu Gln Arg Cys Lys His Pro Phe Leu Thr Glu Leu 245 250 Lys Tyr Ser Phe Gln Thr Asn Asp Arg Leu Cys Phe Val Met Glu Phe 265 Ala Ile Gly Gly Asp Leu Tyr Tyr His Leu Asn Arg Glu Val Gln Met 280 285 Asn Lys Glu Gly Phe Ser Glu Pro Arg Ala Arg Phe Tyr Gly Ser Glu 295 300 Ile Val Leu Ala Leu Gly Tyr Leu His Ala Asn Ser Ile Val Tyr Arg 310 315 Asp Leu Lys Leu Glu Asn Leu Leu Asp Lys Asp Gly His Ile Lys 325 330 Ile Ala Asp Phe Gly Leu Cys Lys Glu Glu Ile Ser Phe Gly Asp Lys 345 -Thr Ser Thr Phe Cys Gly Thr Pro Glu Tyr Leu Ala Pro Glu Val Leu 360 Asp Asp His Asp Tyr Gly Arg Cys Val Asp Trp Trp Gly Val Gly Val 375 380 Val Met Tyr Glu Met Met Cys Gly Arg Leu Pro Phe Tyr Ser Lys Asp 390 395 His Asn Lys Leu Phe Glu Leu Ile Met Ala Gly Asp Leu Arg Phe Pro 405 410 Ser Lys Leu Ser Gln Glu Ala Arg Thr Leu Leu Thr Gly Leu Leu Val 425 Lys Asp Pro Thr Gln Arg Leu Gly Gly Pro Glu Asp Ala Leu Glu 440 Ile Cys Arg Ala Asp Phe Phe Arg Thr Val Asp Trp Glu Ala Thr Tyr

```
Arg Lys Glu Ile Glu Pro Pro Tyr Lys Pro Asn Val Gln Ser Glu Thr
465
                    470
                                        475
Asp Thr Ser Tyr Phe Asp Asn Glu Phe Thr Ser Gln Pro Val Gln Leu
                485
                                    490
Thr Pro Pro Ser Arg Ser Gly Ala Leu Ala Thr Val Asp Glu Gln Glu
                                505
            500
Glu Met Gln Ser Asn Phe Thr Gln Phe Ser Phe His Asn Val Met Gly
                            520
Ser Ile Asn Arg Ile His Glu Ala Ser Glu Asp Asn Glu Asp Tyr Asp
                        535
Met Gly
545
<210> 156
<211> 483
<212> PRT
<213> Caenorhabditis elegans
<400> 156
Met Ser Thr Glu Asn Ala His Leu Gln Lys Glu Asp Ile Val Ile Glu
                                   10
Ser Trp Leu His Lys Lys Gly Glu His Ile Arg Asn Trp Arg Pro Arg
                                25
Tyr Phe Ile Leu Phe Arg Asp Gly Thr Leu Leu Gly Phe Arg Ser Lys
                            40
                                                45
Pro Lys Glu Asp Gln Pro Leu Pro Glu Pro Leu Asn Asn Phe Met Ile
                       55
                                            60
Arg Asp Ala Ala Thr Val Cys Leu Asp Lys Pro Arg Pro Asn Met Phe
                   70
                                        75
Ile Val Arg Cys Leu Gln Trp Thr Thr Val Ile Glu Arg Thr Phe Tyr
               85
                                    90
Ala Asp Ser Ala Asp Phe Arg Gln Met Trp Ile Glu Ala Ile Gln Ala
           100
                               105
Val Ser Ser His Asn Arg Leu Lys Glu Asn Ala Gly Asn Thr Ser Met
        115
                           120
                                                125
Gln Glu Glu Asp Thr Asn Gly Asn Pro Ser Gly Glu Ser Asp Val Asn
                       135
                                            140
Met Asp Ala Thr Ser Thr Arg Ser Asp Asn Asp Phe Glu Ser Thr Val
                   150
                                        155
Met Asn Ile Asp Glu Pro Glu Glu Val Pro Arg Lys Asn Thr Val Thr
                                    170
               165
                                                        175
Met Asp Asp Phe Asp Phe Leu Lys Val Leu Gly Gln Gly Thr Phe Gly
           180
                               185
Lys Val Ile Leu Cys Arg Glu Lys Ser Ser Asp Lys Leu Tyr Ala Ile
                            200
                                                205
Lys Ile Ile Arg Lys Glu Met Val Val Asp Arg Ser Glu Val Ala His
                        215
                                            220
Thr Leu Thr Glu Asn Arg Val Leu Tyr Ala Cys Val His Pro Phe Leu
                   230
                                        235
Thr Leu Leu Lys Tyr Ser Phe Gln Ala Gln Tyr His Ile Cys Phe Val
                245
                                    250
Met Glu Phe Ala Asn Gly Gly Glu Leu Phe Thr His Leu Gln Arg Cys
                                265
                                                    270
Lys Thr Phe Ser Glu Ala Arg Thr Arg Phe Tyr Gly Ser Glu Ile Ile
                            280
                                                285
Leu Ala Leu Gly Tyr Leu His His Arg Asn Ile Val Tyr Arg Asp Met
                        295
                                            300
Lys Leu Glu Asn Leu Leu Asp Arg Asp Gly His Ile Lys Ile Thr
```

```
305
                    310
                                        315
Asp Phe Gly Leu Cys Lys Glu Glu Ile Lys Tyr Gly Asp Lys Thr Ser
               325
                                   330
Thr Phe Cys Gly Thr Pro Glu Tyr Leu Ala Pro Glu Val Ile Glu Asp
            340
                                345
Ile Asp Tyr Asp Arg Ser Val Asp Trp Trp Gly Val Gly Val Val Met
        355
                            360
Tyr Glu Met Met Cys Gly Arg Leu Pro Phe Ser Ala Lys Glu Asn Gly
                        375
                                            380
Lys Leu Phe Glu Leu Ile Thr Thr Cys Asp Leu Lys Phe Pro Asn Arg
                    390
                                        395
Leu Ser Pro Glu Ala Val Thr Leu Leu Ser Gly Leu Leu Glu Arg Val
                405
                                    410
                                                        415
Pro Ala Lys Arg Leu Gly Ala Gly Pro Asp Asp Ala Arg Glu Val Ser
            420
                                425
Arg Ala Glu Phe Phe Lys Asp Val Asp Trp Glu Ala Thr Leu Arg Lys
                            440
                                                445
Glu Val Glu Pro Pro Phe Lys Pro Asn Val Met Ser Glu Thr Asp Thr
                       455
                                            460
Ser Phe Phe Asp Arg Val Arg Tyr Val Ser Ile Leu Leu Lys Val Ser
                   470
                                        475
Glu Ala Ile
```

<210> 157 <211> 480 <212> PRT

<213> Homo sapiens

<400> 157 Met Ser Asp Val Ala Ile Val Lys Glu Gly Trp Leu His Lys Arg Gly Glu Tyr Ile Lys Thr Trp Arg Pro Arg Tyr Phe Leu Leu Lys Asn Asp Gly Thr Phe Ile Gly Tyr Lys Glu Arg Pro Gln Val Asp Val Gln Arg Glu Ala Pro Leu Asn Asn Phe Ser Val Ala Gln Cys Gln Leu Met Lys Thr Glu Arg Pro Arg Pro Asn Thr Phe Ile Ile Arg Cys Leu Gln Trp

Thr Thr Val Ile Glu Arg Thr Phe His Val Glu Thr Pro Glu Glu Arg Glu Glu Trp Thr Thr Ala Ile Gln Thr Val Ala Asp Gly Leu Lys Lys Gln Glu Glu Glu Met Asp Phe Arg Ser Gly Ser Pro Ser Asp Asn Ser Gly Ala Glu Glu Met Glu Val Ser Leu Ala Lys Pro Lys His Arg Val Thr Met Asn Glu Phe Glu Tyr Leu Lys Leu Leu Gly Lys Gly Thr Phe Gly Lys Val Ile Leu Val Lys Glu Lys Ala Thr Gly Arg Tyr Tyr Ala Met Lys Ile Leu Lys Lys Glu Val Ile Val Ala Lys Asp Glu Val Ala His Thr Leu Thr Glu Asn Arg Val Leu Gln Asn Ser Arg His Pro Phe Leu Thr Ala Leu Lys Tyr Ser Phe Gln Thr His Asp Arg Leu Cys

```
ddssssfact cydssfact gastegigc tigaagiaag ctigcccatt titticggaa
II40
        वेत्तेषष्ठतेषे वेषष्ठाद्वात् वेषष्ठव्तेत्वेत् वष्ठाव्यत्वात् व्यवेष्ठवेष्ठ वेषष्ठष्ठवेत्त्व
1080
        adaacticca acgacticat gittcticag agtaigggog aaggagccia cagccaggit
1050
        fiatococad toaccyccya agaicteata yotaaaayca ttaaagaayy atytocyaay
096
006
        casatitita tittocagac gotoaactta acaccaacag caagigaato ggagaacago
018
        titcacatit tacaactati ciaggcaaa aigaaaaaa aaaactigia gaataatiti
        atattaatta aatatgaatt togaaatatg aattttggtt gaottooatt atgtttttt
081
        tttcacctgc ctaagatcgt tttagcataa atatgtagat gaccgagagt atacaattaa
150
        acyclctaa actigaatti gaaaattig agtigatgot ttaatataaa agtittgagg
099
        tittaggett tittaaata aatgiitggg eeggaacaet taacegaata geatgaiga
009
        stygettety tttgeetact tetagettga acattetaay geteegage gaaaaatt
015
08Þ
        cradiccear ificadaecd ficeaetiff acctacaata caaaatigge eggeageti
        aagttaaata gaaatattt taaaatattt tttttgtot aggaaaatt gataaagcac
450
        adadccaatt caaccggaaa actettttt atagggaaaa cgttttgcca cgtagcagat
390
        aatacactit tigaacctaa aacctagatt titigigitc taaatictit igigaatigg
300
        redddidese fiateaact agaceegit titagaeeet etiteaage gggaeige
240
08T
        двядсадсас свясадідад двясівдії сівдясдвяс ягоддвядс ддсігвявді
        acaccaacta acacgtogot cgacaccaca actactaaca atgacacgac atcggatcgt
150
        cataaaaatc cagtaaatgg taaaattttc aatttcagat ccatctcgat ggaggatctc
09
                                                                 <4005>
```

<213> Caenorhabditis elegans

AND <SIS>

<511> 6250

<210> 128

```
084
                                                             99 b
                    SLD
                                         015
Arg Arg Pro His Phe Pro Gln Phe Ser Tyr Ser Ala Ser Ser Thr Ala
                09 Þ
                                     95£
Ile Thr Pro Pro Asp Gln Asp Ser Met Glu Cys Val Asp Ser Glu
            GFF
                                Obb
Lyr Asp Thr Arg Tyr Phe Asp Glu Glu Phe Thr Ala Gln Met Ile Thr
        430
                            452
                                                 02 b
IAL GIM PAS PAS TEW SER PRO PRO PAS PRO GIM VAL TAR GEW GIM
    SIB
                        OTF
                                             SOF
Glu Ile Met Gln His Arg Phe Phe Ala Gly Ile Val Trp Gln His Val
00Þ
                    362
                                         360
rha rha yab bio rha cin yid ren ciy ciy cer ciu yap ala rya
                380
                                     375
                                                         370
bro Arg Thr Leu Gly Pro Glu Ala Ser Leu Leu Ser Gly Leu Leu
                                09€
            365
                                                     322
Asp His Glu Lys Leu Phe Glu Leu Ile Leu Met Glu Glu Ile Arg Phe
                            342
                                                 340
Val Val Met Tyr Glu Met Met Cys Gly Arg Leu Pro Phe Tyr Asn Gln
    332
                        330
                                             325
ren ejn yab yau yab Iyr ely Arg Ala Val Asp Trp Gly Leu Gly
350
                    312
                                         310
LUR MET LYS Thr Phe Cys Gly Thr Pro Glu Tyr Leu Ala Pro Glu Val
                300
                                     562
                                                         067
rys Ile Thr Asp Phe Gly Leu Cys Lys Glu Gly Ile Lys Asp Gly Ala
            582
                                280
                                                     SLZ
yrd ysb ren rha ren cjn ysu ren Wet ren ysb rha ysb clh His Ile
        01.2
                            265
                                                 092
Ile Val Ser Ala Leu Asp Tyr Leu His Ser Glu Lys Asn Val Val Tyr
    255
                                             545
                        220
yrd cjn yrd Ast bye Ser cjn ysb bye yjs bye yrd Iyr cjy yjs cjn
240
                    235
                                         230
Phe Val Met Glu Tyr Ala Asn Gly Glu Leu Phe His Leu Ser
```

089₽ geatgatic titigeatet actgateaa aatigatica aateaatiaa titictiiga titicatity cactyaccat cicticatit goactyacca actiticati tycaaticty **4**620 cadifigoac igaccaccic itcatiigoa cigaccacci citcatiigo acigaccaac 0955 tgattaacaa tgaccaaaag atttgaactg acaaagtgca aatttgcacc gaccaaaaa 00SF dosfiasad tittacotty cactdacca aattatty aactattaat tattigatto 0110 4380 rridddagar dardcradcd carcacadcc arcaacdrda drridaadca rrirrrcrr croracati gggcctgtcg agcogggact tgatgatcgt gccttgttcc gtttgatgaa 4350 **45**00 tateaageca ceagteetge acgeetaeat tecageeaea tttggegage eggagtaeta cadicaagaa citaiggcic acaagiiit igaaaacgii gacigggiga acatigcaaa **4**500 rrasadac gaatacgtaa tittoaact acaggigogo gaccogagia cocgiatoac 0 F T F addarffcca gaggaaggt cggaaaft teggraggt gacatgaaac 4080 adcodicase cadiaceate titigaaag aatecaggag tiggatitet egiteceaga 4050 cdacatttgg gaattgggat gtatcctttt ccagtgtcta gccggacagc caccattcag 0968 adsocseed sadeteeds tettegas atgessatt taacagtigg attecagase 3900 उटउपट्राह्त ह्रितेष्वप्रदाव ह्रिट्रपट्रे व्यटट्रितेष्ठेष ह्रिट्रप्रेट्रिव ट्रिकेप्रवाहे 3840 fffcggaaa fffgactgaa acaattfffg ccagttccag aagagaacac tgctcgacgt 3780 dercreade rectadered cedecaces creatrer recaderaed dedadreedd 3120 3660 carreadeda rerecesera resesadada aerrrsedas racassresa aessaereae scsstdtdct cstccsdssa dacggtcsca ttctcstcsc agattttgga agtgcccagg 3600 recreacedd aerdeaatte ctacaegaea aeaaaattgt geacagagae atgaageegg 3240 adicacidia ccattitada teaticaaca tacteacete aaaatictit gecteggaaa 3480 3450 raaacctcaa ttaatattca gatttcgtga tcggacttgt tgaaaatggt gatcttggcg ccatcaaaat ccacttgtga tcattttatt ccaataaaaa cgtcaactta aaaaaaaaat 3360 3300 attiteacya ccayyctaya attiyiqayi titticcayc yccaayytto tittoiyaac tettaacata cetgteacaa gaatgeggtg gteateegtt tgteacaeag etetacaeae 3540 recadadre gracereae egecateaa aaatggaege aateattege gagaagata 3180 refreceddr afrecdardr ededaadigg eaecadaige gardriegee greaagige 3150 tigtagacyc acctaaaaa titcaaaaa ccaaaaaca aycticcayt aaaacoctaa 3060 3000 tataattago aaaacaataa gtaacattto tgaaaaatta gaacotttoo ogcattgtat aatitaccyc aaacicitca actyaayoca ciatiycaca tiaaciyica aaaticiyya 2940 rraradrara racadacass racadasses rarassas serrradas 2880 csciddccss satigagett geacigaceg agitiagega ceataietti titgietaat 2820 sacidiaagi igaacgaati tacaataaa aacacagcig cacigaccaa aaaacaatta 5160 2700 titateaact gacactgata atatiticig coteatata aaaaatatic otetageaaa 5640 reaacgaatt ticagettee aaattttggt egtitttgga tettittaca aaaaaaatat ttgaatagca aatcttgaaa acgtaaaaac aataattatt ttctatatct gtaaatattt 2280 actagitice geceeece cetataeata igaigeacae tiaaaaigie caagiggigi 2220 agtiglaaac actitgatag ttaaaatgat tgttigtagt gatcagaagc agaaaatctg 2460 agtgaatgga aatttottga ctaaatoogt ggaaaattat ctagttttgt ttttoagata 2400 2340 ttottaaaat agtogaottg aaataatttt togttatta toaatooaat gagttgaaaa aatttatgto gttggaactt ccaatttgga agtacagttt tttggaaatt aaatttttga 2280 rrdsorrrcd dededeeddd rrfdrecror decrerdrer eecroeedee deerdreddd 2220 defitadace etgaggeata titectacae etagaacaa taccgiaatg aatetitaca 5160 cardarrarr adarrosics satigacade geacaggas tigatitiga acquarate 2100 2040 cerdiacity itgaaggaa aaacgagacg ittgigidia itgggggggg gtaatgiaac dedretetta cacaaaataa gecaegegte tageactate aacategea aacagetata 1980 gaagaagatg cagaaaaga cgacatogic atagaatigi ctacacaaac ctagigitci 1920 cccctgtcca atctctttc tccacaacac tttaatctca tttcgcatgg agaagagaaa 098T readddrfac acdaetdfdd daartdaact cgeactatgt aggeceatte atgttgfetet 1800 JJ40 ceeeeetcar ecceedecad ecrecrarer rrrradicad eeeeercadc ceerrraca daccdraaca ctiticcaat ggcgtataca attigaatit agcaacaaaa caaaaaaaa 089T tittactyta ayttatcatc attitggcac cgaaaggtit tittaggtaa tittaccact T 620 ataticcaac atigtaggaa tictagaati gotitagati tototitgit ticcaatoti 09ST trictagics tigaatitis accticosat tittaigsts taacigigit teaaatacte 1200 tadcactasa atgagcasct atcataagaa attagaaaat ttggaaaatt ggtttatttt 1440 ticaaatoti titaaaggit tagtacggic attaaaaaa atatttaaaa atcatotica 1380 tyayttta togattta tagcttttt tacttatata tattcaaaat gtatgtgtt 1350 attigcagit titgccacaa atciaictig acacaataia coicaciati agitaaaigc **T**560 catoddfdat tofftoffgg caattcaact gatagtactg gtattaccta gccgcaaaaa 1500

bye yfa Ser Glu Ile Leu Thr Gly Leu Gln Phe Leu His Asp Asn Lys SLT OLT 59 T GJn Ser Leu Cys His Phe Gly Ser Phe Asp Met Leu Thr Ser Lys Phe OST 09T **T22** STT yja yıd ije ili bie val ile Giy Leu val Giu Asn Giy Asp Leu Giy IdO **332** UET CJY GLY His Pro Phe Val Thr Gln Leu Tyr His Phe His Asp Gln 150 SII Ala ile ile Arg Glu Lys Asn ile Leu Thr Tyr Leu Ser Gln Glu Cys OII SOT OOT Val Lys Val Leu Gln Lys Ser Tyr Leu Asn Arg His Gln Lys Met Asp 06 Ser Gln Val Phe Arg Cys Arg Glu Val Ala Met Phe Ala 08 9 INT SET ASD ASP PRE PRE LOG GIN SET MET GLY GLU GLY ALA TYR 99 09 YIS GIU ASP Leu Ile Ala Lys Ser Ile Lys Glu Gly Cys Pro Lys Arg 0₽ Teu Thr Pro Thr Ala Ser Glu Ser Glu Asn Ser Pro Val Thr 30 yau yau yab Ipr Jer Asp Arg Glu Ala Ala Pro Thr Leu Asn ST OI WET GIN WSP LEU Thr Pro Thr Asn Thr Ser Leu Asp Thr Thr Thr 6ST <000>>

<213> Caenorhabditis elegans

<212> PRT

<211>

<210> 129

rrrdcdrcc 9229 ddcfcacctc coffctactc cocacaaat caccatcaaa caaatcacac fttfgtatca 6240 0819 reandcode dosedrande esquadorir caardosaar agacaagaag rogoorigaa sestderdcd rdsscedesd dcdcrdcdcc dcssscesds sesddedded sesseddcdc 9170 rrcaatactt cttttctgtt caaaaatga tcaacttgct gaaaaaaat tttttgtagg 0909 aagcatttt ggaccaccac gcaaatctta ttattatgga ccacccaaac ttagaacacc 0009 attttggacc accctgtata attatggacc accatgtaca cttatagacc acccagtaac 0165 2880 ttatgatgtt tagacggttt aaatttttg atgatttaaa tttttaggg gtggtctata taaaggitot ttatiticca igcaacicta aaatoticco giatatiti tiggaaagic 2850 agattotgga aaattitoaa aaaagataa tototaaaca aaactaaatt caaaatgito 0915 aaagaatata gaatatttgt gttcaacttt tcttgtcaaa atattttttt tggacaatct 0049 aaattotytt otoaaaatty gattittaca gayottytt oyagatttoa taatoottoa 0199 casaaaaaa ttgagaaatg tctgaaaaa tttggagtgt gacagttttc tgaattttga 0899 tdccaatttt cattagactt tagagcctat tgctatttg tggacaggtt taaacatttt 2220 fractggaag goccocctca ctgagtitoc agcaagtica gagtitita tiggaatiti 0979 cdsfdcdfds cddsscsfff ddcsdcsfff sfddsssds ssddfccsds ssddfsfdss 2400 addotatosa tgatotogo aagoggact ogtgactat ogaaagact ttaactotg 2340 freageceaa cegegietae tactigitig atcicgaaa gaaageagat gagiggigta 2280 cagaataatc atagctgtct atctcattat agtactcaat gaatctgaaa atttcaaatt 2220 acgccgtgca tgcaggtgga gctaaaaac tcgggaactt tctttataca tacggtaggt 0915 ddsccdcsfc foffgtscsf fgatgfgcog safoffgtgc fcsaaggaga ggtaccatgg 0019 aderettigg asasgasges agaitgitt geesgeege gaatgiteet gitgaeegaa 2040 gageacgig teaaaacce attecaeate tteaceaca actegeteat tttgaaaca 0867 faradacdcd ccaaticdda adccdaaad aaccdcdccd cacdidcdca gaagcicdaa 4920 ficggaaatt gcaccgigag igattigcac ctaatiggit attittaata atcattaaat 0987 tactaaccc ttagaaatt tcagacctc taacgiggaa catcgcggag acccattgt 4800 cagtactatg cottattcaa ggagatgotg atotgaaaat totcaatagt tgataaaaat 01/10

089 Wet Gln Met Asp Lys Lys Ser Pro 029 ST9 Glu Glu Lys Lys Ala Leu Lys Ala Glu Gln Val Ser Lys Lys Leu Ser Lys Glu Met Met Arg Glu Gln Lys Ala Leu Arg Arg Lys Gln Glu Lys Wer yrd ysb Gly Thr Phe Gly Ser 1le Tyr Gly Lys Lys Ser Arg Val Arg Lys Arg Tyr Ser Val Thr 11e Glu Lys Thr Phe Asn Ser Ala 099 Phe Asp Leu Glu Lys Ala Asp Glu Trp Cys Lys Ala Ile Asn Asp Asn Ser Gly Thr Phe Phe Ile His Thr Pro Asn Arg Val Tyr Leu Leu Lys Gly Glu Val Pro Trp Thr Pro Cys Met Gln Val Glu Leu Lys ren Ibr Glu Gly Pro His Leu Leu Tyr 1le Asp Val Pro Asn Leu Val Ill Leu Glu Lys Lys Arg Gly Leu Phe Ala Arg Arg Met Phe Leu yau bro Phe His Ile Phe Thr Asn Asn Ser Leu Ile Leu Lys Gln Gly rks yeu yrd yla yla brd hla Gln Glu Glu Brg Val Lys Asp Pro Phe Val Ser Glu Ile Ala Pro Arg Ala Asn Ser Glu Ala Glu Ala Ser Ala Ser Gln Pro Ser Thr Pro Ser Asn Val Glu His Arg Gly Gly Leu Asp Asp Arg Ala Leu Phe Arg Leu Met Asn Leu Gly Asn Asp Lyr Phe Gly Glu Fro Glu Tyr Tyr Ser Asn 1le Gly Pro Val Glu Pro Asn Ile Ala Asn Ile Lys Pro Pro Val Leu His Ala Tyr Ile Pro Ala Ser Gln Glu Leu Met Ala His Lys Phe Phe Glu Asn Val Asp Trp Val 345 Clu lle ile Ala Lys Ile Leu Val Arg Asp Pro Ser Thr Arg Ile Thr Cln Glu Leu Asp Phe Ser Phe Pro Glu Gly Phe Pro Glu Glu Ala Ser Cln Pro Pro Phe Arg Ala Val Asn Gln Tyr His Leu Leu Lys Arg Ile Thr Asp ile Trp Gly Leu Gly Cys ile Leu Phe Gln Cys Leu Ala Gly Leu Tyr Val Ser Pro Glu Met Leu Ala Asp Gly Asp Val Gly Pro Gln 592 CIN CIN CIN CIN YOU INK YIS YED YED INK INK BUG AST CIN INK YIS Arg Ser Ser Asp Ser Gly Ser Pro Pro Thr Arg Phe Tyr Ser Asp 232 230 ren eju ren ser eju ejn ejn be ihr kap Ala Asn Gin Ala Ser Ser 512 Gly His Ile Leu Ile Thr Asp Phe Gly Ser Ala Gln Ala Phe Gly Gly 200 Ite Val His Arg Asp Met Lys Pro Asp Asn Val Leu Ile Gln Lys Asp 06 T **381** 180

yau yab yia Ser Ala Ser Gin Pro Ser Thr Phe Arg Pro Ser Asn Val OTF S01 ein Pro Gly Leu Asp Asp Arg Ala Leu Phe Arg Leu Met Asn Leu Gly 390 OOF 368 LIO WIS The Pre Gly Glu Pro Glu Tyr Tyr Ser Asn Ile Gly Pro Val SLE Trp Val Asn Ile Ala Asn Ile Lys Pro Pro Val Leu His Ala Tyr Ile 365 360 322 Ile Thr Ser Gln Glu Leu Met Ala His Lys Phe Phe Glu Asn Val Asp 340 345 Ala Ser Glu Ile Ile Ala Lys Ile Leu Val Arg Asp Pro Ser Thr Arg 330 yrd ile Gin Geu Asp Phe Ser Phe Pro Giu Giy Phe Pro Giu Giu 312 310 Ala Gly Gln Pro Pre Arg Ala Val Asn Gln Tyr His Leu Leu Lys 262 Pro Gln Thr Asp lle Trp Gly Leu Gly Cys lle Leu Phe Gln Cys Leu 280 285 Thr Ala Leu Tyr Val Ser Pro Glu Met Leu Ala Asp Gly Asp Val Gly 597 Glu Glu Val Pro Glu Glu Asn Thr Ala Arg Arg Thr Thr Phe Val Gly 545 720 yıd ger ger yab ger gji ger bro Pro Thr Arg Phe Tyr Ser Asp 232 230 ren eju ren set ejn ejn ejn bye Thr Asp Ala Asn ein Ala Ser ser 512 Gly His 11e Leu 11e Thr Asp Phe Gly Ser Ala Gln Ala Phe Gly Gly 200 202 lle Val His Arg Asp Met Lys Pro Asp Asn Val Leu Ile Gln Lys Asp **T82** Phe Ala Ser Glu Ile Leu Thr Gly Leu Gln Phe Leu His Asp Asn Lys 59 T OLT Glu Ser Leu Cys His Phe Gly Ser Phe Asp Met Leu Thr Ser Lys Phe 122 120 yra yrd 116 Ilx bye nal 11e Gly Leu Val Glu Asn Gly Asp Leu Gly 332 GLY GLY His Pro Phe Val Thr Gln Leu Tyr His Phe His Asp Gln 150 yia ile ile Arg Glu Lys Asn ile Leu Thr Tyr Leu Ser Gln Glu Cys SOT Val Lys Val Leu Gln Lys Ser Tyr Leu Asn Arg His Gln Lys Met Asp 06 Ser Gln Val Phe Arg Cys Arg Glu Val Ala Thr Asp Ala Met Phe Ala SLThr Ser Asn Asp Phe Met Phe Leu Gln Ser Met Gly Glu Gly Ala Tyr 99 yra cin ysp Leu ile Ala Lys Ser ile Lys Glu Gly Cys Pro Lys Arg OΦ ren Ibr Pro Thr Ala Ser Glu Ser Glu Asn Ser Leu Ser Pro Val Thr Asn Asn Asp Thr Ser Asp Arg Glu Ala Ala Pro Thr Leu Asn 0 T Met Glu Asp Leu Thr Pro Thr Asn Thr Ser Leu Asp Thr Thr Thr Thr 091 <000>

<213> Caenorhabditis elegans

<212> PRT

^{989 &}lt;117>

<510> 100

<511> 2d <510> 103

<211> 28 <211> 28 <211> PRT <213> Homo sapiens

> <211> 54 <211> 54 <213> Homo sapiens

089 Lys Lys Leu Ser Met Gln Met Asp Lys Ser Pro 920 919 rks eju ejn rks ejn ejn rks rks vja ren rks vja ejn ejn Nal Ser 009 TYS LYS Ser Arg Lys Glu Met Met Arg Glu Gln Lys Ala Leu Arg Arg 282 bye yau ser yjs Wef yrd yab Cjh lyr bye Cjh ser ije lhr Cjh rha 019 yfs lje yau yab yat yrg Lyr Ser yal Thr lle Glu Lyr Thr 222 055 Val Tyr Leu Phe Asp Leu Glu Lys Lys Ala Asp Glu Trp Cys Lys 532 Val Glu Leu Lys Asn Ser Gly Thr Phe Phe 11e His Thr Pro Asn Arg 250 bro yeu ren val Leu Lys Gly Glu val Pro Trp Thr Pro Cys Met Gln 909 Arg Met Phe Leu Leu Thr Glu Gly Pro His Leu Leu Tyr 1le Asp Val 06ħ ren ris eju eji Lit ren ejn ris ris yta eji ren bye yja yta yta SLF Cln Arg Val Lys Asn Pro Phe His 11e Phe Thr Asn Asn Ser Leu 11e SSF Ser Glu Ala Glu Lys Asn Arg Ala Ala Ala Gln Lys Leu Glu Glu 0 Þ Þ Glu His Arg Gly Asp Pro Phe Val Ser Glu Ile Ala Pro Arg Ala Asn

-82-CIN WIS Pro Gin Val Val Glu Ile Asp Pro Asp Phe Glu Pro Leu Pro L9T <00b> <213> Homo sapiens <212> PRT <211> 45 L9T <0TZ> 20 pro Arg Pro Glu Ser Pro 0Τ ς yrs yeb bro yeb bhe Gin bro Arg Per Cys Thr Trp Peo Leu 991 <001> <213> Homo sapiens <212> PRT <211> 22 991 <017> ΟĐ 32 egu pro ser elu pro pro elu Val elu Pro 3.0 52 yrd bro yrd Ser Cys Thr Trp Pro Leo Ard Pro Glu 1le Ala Asn 0 T Lys Ala Ala Ala Ile Ile Asp Leu Asp Pro Asp Phe Glu Pro Gln Ser S9T <000> <213> Homo sapiens <212> PRT <211> 42 <510> 165 OΤ τ Ser Phe Arg Pro Arg Thr Gin Ser Asn Leu Ser Ser Ser Ser Ser Phe Arg Pro Giy Ser Ser Ser ₱9T <00₱> <213> Caenorhabditis elegans <212> PRT <211> 17 <510> 164 9 Gln Asp Asp Leu Gly Glu ςĐ OΦ 35 Ser Asn Ala Ser Thr Ile Ser Gly Arg Leu Ser Pro Ile Met Thr Glu 25 Ser Asn Asp Asp Pap Asp Asp Trp Ser Thr Phe Arg Pro Arg Thr Ser 0Τ Ser Pro Gly Ser Gln Phe Ser Lys Trp Pro Ala Ser Pro Gly Ser His

> <213> Homo sapiens <212> PRT

E9T <00b>

bye yra yrd yrd ren ren Lyr Clu Cly Pro His Leu Tyr Asp Asn ren ejn ejn yau bro His bhe yan Leu lle Leu Lys Gly Lys Gly Leu <213> Mus musculus or Homo sapiens or C elegans <212> PRT **LP <III>** <510> 111 SL CIn Leu Arg Pro Glu Ala Lys Asn Phe Lys Thr Phe Phe Val His Thr Agy Asp Pro Val Ash Lys Val Leu Lys Gly Glu Ile Pro Trp Ser Gln Ala Arg Arg Gln Leu Leu Thr Glu Gly Pro His Leu Tyr Tyr 52 yau ren 11e ren ria wet eji bro val asp iya arg Lis Giy Leu Phe 0 T ren ejn rys ein Ala Gly ely Asn Pro Trp His ein Phe Val elu Asn <213> Mus musculus or Homo sapiens <212> PRT <211> 80 <210> 110 Thr Pro Val Asp Glu Pro Pro Arg Thr Trp Pro Arg Pro 69T <00b> <213> Caenorhabditis elegans or Homo sapiens <SIS> PRT <211> 14 <510> 169 ren ejn bro bro ben yau ser ser bro bro Ile Pro Arg Asp Arg Cys Asn Thr Trp Pro Met Arg Pro Gln OT. Lyr Phe Met Asn Thr Pro Asp Ash Wet Met Asn Asp Met Glu 89T <00b> <213> Caenorhabditis elegans <212> PRT <211> 41 <510> 168 0 Þ 35 Ser Asn Ser Ala Thr Ser Ser Pro Ala Pro 52 07 Yrd bro Yrd Ser Cys Thr Trp Pro Leu Pro Ard Pro Glu Phe Ser Gln

SI

```
35
<210> 172
<211> 80
<212> PRT
<213> Caenorhabditis elegans
<400> 172
Leu Glu Glu Gln Arg Val Lys Asn Pro Phe His Ile Phe Thr Asn Asn
Ser Leu Ile Leu Lys Gln Gly Tyr Leu Glu Lys Lys Arg Gly Leu Phe
                                25
                                                     30
Ala Arg Arg Arg Met Phe Leu Leu Thr Glu Gly Pro His Leu Leu Tyr
Ile Asp Val Pro Asn Leu Val Leu Lys Gly Glu Val Pro Trp Thr Pro
Cys Met Gln Val Glu Leu Lys Asn Ser Gly Thr Phe Phe Ile His Thr
                                        75
<210> 173
<211> 113
<212> PRT
<213> Mus musculus or Homo sapiens
<400> 173
Ser Asp Leu Trp Ala Leu Gly Cys Ile Ile Tyr Gln Leu Val Ala Gly
Leu Pro Pro Phe Arg Ala Gly Asn Glu Tyr Leu Ile Phe Gln Lys Ile
Ile Lys Leu Glu Tyr Asp Phe Pro Glu Lys Phe Pro Lys Ala Arg
Asp Leu Val Glu Lys Leu Leu Val Leu Asp Ala Thr Lys Arg Leu Gly
Cys Glu Glu Met Glu Gly Tyr Gly Pro Leu Lys Ala His Pro Phe Phe
65
                    70
                                        75
Glu Ser Val Thr Trp Glu Asn Leu His Gln Gln Thr Pro Pro Lys Leu
                85
Thr Ala Tyr Leu Pro Ala Met Ser Glu Asp Asp Glu Asp Cys Tyr Gly
                                105
Asn
<210> 174
<211> 48
<212> PRT
<213> Mus musculus or Homo sapiens or C elegans
<400> 174
Asp Trp Leu Gly Cys Ile Gln Ala Gly Pro Pro Phe Arg Ala Asn Tyr
1
                                    10
Ile Leu Phe Pro Glu Phe Ala Lys Leu Val Leu Glu Pro Leu Ala His
                                25
Phe Phe Glu Val Trp Asn Pro Pro Leu Ala Tyr Pro Ala Glu Tyr Asn
```

Val Leu Lys Gly Glu Pro Trp Glu Lys Asn Thr Phe Phe His Thr

```
<210> 175
<211> 122
<212> PRT
<213> Caenorhabditis elegans
<400> 175
Thr Asp Ile Trp Gly Leu Gly Cys Ile Leu Phe Gln Cys Leu Ala Gly
                                    10
Gln Pro Pro Phe Arg Ala Val Asn Gln Tyr His Leu Leu Lys Arg Ile
                                25
                                                     30
Gln Glu Leu Asp Phe Ser Phe Pro Glu Gly Phe Pro Glu Glu Ala Ser
                            40
                                                 45
Glu Ile Ile Ala Lys Ile Leu Val Gly His Glu Thr Leu Lys Thr Glu
                        55
Tyr Val Ile Phe Asn Leu Gln Val Arg Asp Pro Ser Thr Arg Ile Thr
65
                    70
                                        75
Ser Gln Glu Leu Met Ala His Lys Phe Phe Glu Asn Val Asp Trp Val
                                    90
                                                         95
                85
Asn Ile Ala Asn Ile Lys Pro Pro Val Leu His Ala Tyr Ile Pro Ala
                                105
Thr Phe Gly Glu Pro Glu Tyr Tyr Ser Asn
        115
<210> 176
<211> 72
<212> PRT
<213> Mus musculus or Homo sapiens
<400> 176
Phe Gly Leu Ser Tyr Ala Lys Asn Gly Glu Leu Leu Lys Tyr Ile Arg
                                    10
Lys Ile Gly Ser Phe Asp Glu Thr Cys Thr Arg Phe Tyr Thr Ala Glu
                                25
Ile Val Ser Ala Leu Glu Tyr Leu His Gly Lys Gly Ile Ile His Arg
                            40
                                                 45
Asp Leu Lys Pro Glu Asn Ile Leu Leu Asn Glu Asp Met His Ile Gln
                        55
Ile Thr Asp Phe Gly Thr Ala Lys
<210> 177
<211> 31
<212> PRT
<213> Mus musculus or Homo sapiens or C elegans
<400> 177
Phe Asn Gly Leu Gly Ser Phe Asp Phe Glu Ile Leu Leu His Ile His
                                    10
Arg Asp Lys Pro Asn Leu Asp His Ile Ile Thr Asp Phe Gly Ala
<210> 178
<211> 72
<212> PRT
```

<213> Caenorhabditis elegans

```
<400> 178
Phe Val Ile Gly Leu Val Glu Asn Gly Asp Leu Gly Glu Ser Leu Cys
                                    10
His Phe Gly Ser Phe Asp Met Leu Thr Ser Lys Phe Phe Ala Ser Glu
                                25
Ile Leu Thr Gly Leu Gln Phe Leu His Asp Asn Lys Ile Val His Arg
                            40
Asp Met Lys Pro Asp Asn Val Leu Ile Gln Lys Asp Gly His Ile Leu
                        55
Ile Thr Asp Phe Gly Ser Ala Gln
                    70
<210> 179
<211> 48
<212> PRT
<213> Mus musculus or Homo sapiens
<400> 179
Tyr Ala Ile Lys Ile Leu Glu Lys Arg His Ile Ile Lys Glu Asn Lys
                                    10
Val Pro Tyr Val Thr Arg Glu Arg Asp Val Met Ser Arg Leu Asp His
            20
                                25
                                                     30
Pro Phe Phe Val Lys Leu Tyr Phe Thr Phe Gln Asp Asp Glu Lys Leu
        35
                            40
<210> 180
<211> 15
<212> PRT
<213> Mus musculus or Homo sapiens or C elegans
Ala Lys Leu Lys Lys Arg Glu Leu His Pro Phe Leu Tyr Phe Asp
<210> 181
<211> 53
<212> PRT
<213> Caenorhabditis elegans
<400> 181
Phe Ala Val Lys Val Leu Gln Lys Ser Tyr Leu Asn Arg His Gln Lys
                                    10
Met Asp Ala Ile Ile Arg Glu Lys Asn Ile Leu Thr Tyr Leu Ser Gln
                                                     30
                                25
Glu Cys Gly Gly His Pro Phe Val Thr Gln Leu Tyr Thr His Phe His
       35
                            40
Asp Gln Ala Arg Ile
    50
<210> 182
<211> 29
<212> PRT
<213> Mus musculus or Homo sapiens
```

<400> 182

```
Pro Asn Arg Thr Tyr Tyr Leu Met Asp Pro Ser Gly Asn Ala His Lys
Trp Cys Arg Lys Ile Gln Glu Val Trp Arg Gln Arg Tyr
<210> 183
<211> 15
<212> PRT
<213> Mus musculus or Homo sapiens or C elegans
<400> 183
Pro Asn Arg Tyr Tyr Leu Asp Ala Trp Cys Ile Val Arg Arg Tyr
                                     10
                                                         15
<210> 184
<211> 28
<212> PRT
<213> Caenorhabditis elegans
<400> 184
Pro Asn Arg Val Tyr Tyr Leu Phe Asp Leu Glu Lys Lys Ala Asp Glu
                                                         15
Trp Cys Lys Ala Ile Asn Asp Val Arg Lys Arg Tyr
            20
<210> 185
<211> 25
<212> PRT
<213> Mus musculus or Homo sapiens
<400> 185
Pro Glu Ser Lys Gln Ala Arg Ala Asn Ser Phe Val Gly Thr Ala Gln
                 5
Tyr Val Ser Pro Glu Leu Leu Thr Glu
            20
<210> 186
<211> 15
<212> PRT
<213> Mus musculus or Homo sapiens or C elegans
<400> 186
Pro Glu Ala Arg Phe Val Gly Thr Ala Tyr Val Ser Pro Glu Leu
                 5
                                     10
<210> 187
<211> 25
<212> PRT
<213> Caenorhabditis elegans
<400> 187
Pro Glu Glu Asn Thr Ala Arg Arg Thr Thr Phe Val Gly Thr Ala Leu
1
Tyr Val Ser Pro Glu Met Leu Ala Asp
```

```
<210> 188
<211> 62
<212> PRT
<213> Caenorhabditis elegans
<400> 188
Lys Arg Thr Ser Asn Asp Phe Met Phe Leu Gln Ser Met Gly Glu Gly
Ala Tyr Ser Gln Val Phe Arg Cys Arg Glu Val Ala Thr Asp Ala Met
                                25
Phe Ala Val Lys Val Leu Gln Lys Ser Tyr Leu Asn Arg His Gln Lys
                            40
                                                 45
Met Asp Ala Ile Ile Arg Glu Lys Asn Ile Leu Thr Tyr Leu
                        55
<210> 189
<211> 21
<212> PRT
<213> Caenorhabditis elegans or Homo sapiens
Lys Asp Phe Phe Gly Glu Gly Ser Val Arg Glu Ala Thr Ala Lys Leu
Lys Lys Arg Glu Leu
<210> 190
<211> 62
<212> PRT
<213> Homo sapiens
<400> 190
Lys Lys Arg Pro Glu Asp Phe Lys Phe Gly Lys Ile Leu Gly Glu Gly
                                    10
Ser Phe Ser Thr Val Val Leu Ala Arg Glu Leu Ala Thr Ser Arg Glu
                                25
Tyr Ala Ile Lys Ile Leu Glu Lys Arg His Ile Ile Lys Glu Asn Lys
                            40
Val Pro Tyr Val Thr Arg Glu Arg Asp Val Met Ser Arg Leu
<210> 191
<211> 90
<212> PRT
<213> Caenorhabditis elegans
<400> 191
His Pro Phe Val Thr Gln Leu Tyr Thr His Phe His Asp Gln Ala Arg
                                    10
Ile Tyr Phe Val Ile Gly Leu Val Glu Asn Gly Asp Leu Gly Glu Ser
            20
                                25
                                                     30
Leu Cys His Phe Gly Ser Phe Asp Met Leu Thr Ser Lys Phe Phe Ala
        35
                            40
```

Ser Glu Ile Leu Thr Gly Leu Gln Phe Leu His Asp Asn Lys Ile Val 55 His Arg Asp Met Lys Pro Asp Asn Val Leu Ile Gln Lys Asp Gly His 70 Ile Leu Ile Thr Asp Phe Gly Ser Ala Gln 85 <210> 192 <211> 39 <212> PRT <213> Caenorhabditis elegans <400> 192 His Pro Phe Leu Tyr Phe Asp Tyr Phe Asn Gly Leu Gly Ser Phe Asp 1 Phe Glu Ile Leu Leu His Ile His Arg Asp Lys Pro Asn Leu Asp His 20 25 Ile Ile Thr Asp Phe Gly Ala 35 <210> 193 <211> 90 <212> PRT <213> Homo sapiens <400> 193 His Pro Phe Phe Val Lys Leu Tyr Phe Thr Phe Gln Asp Asp Glu Lys 1 Leu Tyr Phe Gly Leu Ser Tyr Ala Lys Asn Gly Glu Leu Leu Lys Tyr 20 25 Ile Arg Lys Ile Gly Ser Phe Asp Glu Thr Cys Thr Arg Phe Tyr Thr 35 40 Ala Glu Ile Val Ser Ala Leu Glu Tyr Leu His Gly Lys Gly Ile Ile 55 His Arg Asp Leu Lys Pro Glu Asn Ile Leu Leu Asn Glu Asp Met His Ile Gln Ile Thr Asp Phe Gly Thr Ala Lys 85 <210> 194 <211> 98 <212> PRT <213> Caenorhabditis elegans <400> 194 Glu Glu Asn Thr Ala Arg Arg Thr Thr Phe Val Gly Thr Ala Leu Tyr 1 Val Ser Pro Glu Met Leu Ala Asp Gly Asp Val Gly Pro Gln Thr Asp Ile Trp Gly Leu Gly Cys Ile Leu Phe Gln Cys Leu Ala Gly Gln Pro 35

Pro Phe Arg Ala Val Asn Gln Tyr His Leu Leu Lys Arg Ile Gln Glu

Leu Asp Phe Ser Phe Pro Glu Gly Phe Pro Glu Glu Ala Ser Glu Ile

Ile Ala Lys Ile Leu Val Arg Asp Pro Ser Thr Arg Ile Thr Ser Gln

Glu Leu

<210> 195 <211> 43

<212> PRT

<213> Caenorhabditis elegans or Homo sapiens

<400> 195

Glu Ala Arg Phe Val Gly Thr Ala Tyr Val Ser Pro Glu Leu Asp Trp 10 1 Leu Gly Cys Ile Gln Ala Gly Pro Pro Phe Arg Ala Asn Tyr Ile Leu 25

Phe Pro Glu Phe Ala Lys Leu Val Asp Arg Glu

<210> 196

<211> 98

<212> PRT

<213> Homo sapiens

<400> 196

Glu Ser Lys Gln Ala Arg Ala Asn Ser Phe Val Gly Thr Ala Gln Tyr 1 10

Val Ser Pro Glu Leu Leu Thr Glu Lys Ser Ala Cys Lys Ser Ser Asp 25

Leu Trp Ala Leu Gly Cys Ile Ile Tyr Gln Leu Val Ala Gly Leu Pro 40

Pro Phe Arg Ala Gly Asn Glu Tyr Leu Ile Phe Gln Lys Ile Ile Lys 55

Leu Glu Tyr Asp Phe Pro Glu Lys Phe Pro Lys Ala Arg Asp Leu 75

Val Glu Lys Leu Leu Val Leu Asp Ala Thr Lys Arg Leu Gly Cys Glu 90

Glu Met

<210> 197

<211> 35

<212> PRT

<213> Caenorhabditis elegans

<400> 197

Leu Met Ala His Lys Phe Phe Glu Asn Val Asp Trp Val Asn Ile Ala 10

Asn Ile Lys Pro Pro Val Leu His Ala Tyr Ile Pro Ala Thr Phe Gly

Glu Pro Glu 35

<210> 198

<211> 17

<212> PRT

<213> Caenorhabditis elegans or Homo sapiens

```
<400> 198
Leu Ala His Phe Phe Glu Val Trp Asn Pro Pro Leu Ala Tyr Pro Ala
Glu
<210> 199
<211> 35
<212> PRT
<213> Homo sapiens
<400> 199
Leu Lys Ala His Pro Phe Phe Glu Ser Val Thr Trp Glu Asn Leu His
                                    10
Gln Gln Thr Pro Pro Lys Leu Thr Ala Tyr Leu Pro Ala Met Ser Glu
                                25
Asp Asp Glu
        35
<210> 200
<211> 104
<212> PRT
<213> Caenorhabditis elegans
<400> 200
Leu Glu Glu Gln Arg Val Lys Asn Pro Phe His Ile Phe Thr Asn Asn
1
                                    10
Ser Leu Ile Leu Lys Gln Gly Tyr Leu Glu Lys Lys Arg Gly Leu Phe
                                25
                                                    30
Ala Arg Arg Met Phe Leu Leu Thr Glu Gly Pro His Leu Leu Tyr
                            40
Ile Asp Val Pro Asn Leu Val Leu Lys Gly Glu Val Pro Trp Thr Pro
                        55
Cys Met Gln Val Glu Leu Lys Asn Ser Gly Thr Phe Phe Ile His Thr
                    70
                                        75
Pro Asn Arg Val Tyr Tyr Leu Phe Asp Leu Glu Lys Lys Ala Asp Glu
                85
Trp Cys Lys Ala Ile Asn Asp Val
            100
<210> 201
<211> 59
<212> PRT
<213> Caenorhabditis elegans or Homo sapiens
<400> 201
Leu Glu Gln Asn Pro His Phe Asn Leu Ile Leu Lys Gly Lys Gly Leu
1
                                                         15
                                    10
Phe Ala Arg Arg Leu Leu Thr Glu Gly Pro His Leu Tyr Asp Asn
                                25
Val Leu Lys Gly Glu Pro Trp Glu Lys Asn Thr Phe Phe His Thr Pro
                            40
Asn Arg Tyr Tyr Leu Asp Ala Trp Cys Ile Val
```

```
<210> 202
<211> 104
<212> PRT
<213> Homo sapiens
<400> 202
Leu Glu Lys Gln Ala Gly Gly Asn Pro Trp His Gln Phe Val Glu Asn
                - 5
                                    10
Asn Leu Ile Leu Lys Met Gly Pro Val Asp Lys Arg Lys Gly Leu Phe
                                25
                                                    30
Ala Arg Arg Arg Gln Leu Leu Thr Glu Gly Pro His Leu Tyr Tyr
                           40
Val Asp Pro Val Asn Lys Val Leu Lys Gly Glu Ile Pro Trp Ser Gln
                       55
                                            60
Glu Leu Arg Pro Glu Ala Lys Asn Phe Lys Thr Phe Phe Val His Thr
                                        75
                    70
Pro Asn Arg Thr Tyr Tyr Leu Met Asp Pro Ser Gly Asn Ala His Lys
               85
                                    90
Trp Cys Arg Lys Ile Gln Glu Val
            100
<210> 203
<211> 45
<212> PRT
<213> Homo sapiens
<400> 203
Lys Leu Glu Asn Leu Met Leu Asp Lys Asp Gly His Ile Lys Ile Thr
                5
                                    10
Asp Phe Gly Leu Cys Lys Glu Gly Ile Lys Asp Gly Ala Thr Met Lys
                                25
Thr Phe Cys Gly Thr Pro Glu Tyr Leu Ala Pro Glu Val
                            40
<210> 204
<211> 36
<212> PRT
<213> Homo sapiens or Caenorhabditis elegans
<400> 204
Lys Leu Glu Asn Leu Leu Asp Lys Asp Gly His Ile Lys Ile Asp Phe
                5
                                    10
Gly Leu Cys Lys Glu Ile Gly Thr Phe Cys Gly Thr Pro Glu Tyr Leu
            20
Ala Pro Glu Val
        35
<210> 205
<211> 45
<212> PRT
<213> Caenorhabditis elegans
<400> 205
Lys Leu Glu Asn Leu Leu Asp Lys Asp Gly His Ile Lys Ile Ala
Asp Phe Gly Leu Cys Lys Glu Glu Ile Ser Phe Gly Asp Lys Thr Ser
```

```
20
                                25
Thr Phe Cys Gly Thr Pro Glu Tyr Leu Ala Pro Glu Val
        35
                            40
<210> 206
<211> 62
<212> PRT
<213> Caenorhabditis elegans
<400> 206
Leu Cys Lys Glu Glu Ile Lys Tyr Gly Asp Lys Thr Ser Thr Phe Cys
Gly Thr Pro Glu Tyr Leu Ala Pro Glu Val Ile Glu Asp Ile Asp Tyr
            20
Asp Arg Ser Val Asp Trp Trp Gly Val Gly Val Val Met Tyr Glu Met
                            40
Met Cys Gly Arg Leu Pro Phe Ser Ala Lys Glu Asn Gly Lys
    50
                        55
<210> 207
<211> 43
<212> PRT
<213> Caenorhabditis elegans or Mus musculus
Leu Cys Lys Glu Ile Gly Thr Phe Cys Gly Thr Pro Glu Tyr Leu Ala
Pro Glu Val Glu Asp Asp Tyr Arg Val Asp Trp Trp Gly Gly Val Val
Met Tyr Glu Met Met Cys Gly Arg Leu Pro Phe
<210> 208
<211> 492
<212> PRT
<213> Caenorhabditis elegans
<400> 208
Met Gly Val Asn Asp His Asp Val Ser Val Pro Leu Gln Glu Val Gln
Ser Arg Thr Val Glu Gly Lys Leu Thr Lys Cys Leu Ala Phe Ser Ala
                                25
Phe Val Ile Thr Leu Ala Ser Phe Gln Phe Gly Tyr His Ile Gly Cys
                            40
Val Asn Ala Pro Gly Gly Leu Ile Thr Glu Trp Ile Ile Gly Ser His
                        55
Lys Asp Leu Phe Asp Lys Glu Leu Ser Arg Glu Asn Ala Asp Leu Ala
                    70
                                        75
Trp Ser Val Ala Val Ser Val Phe Ala Val Gly Gly Met Ile Gly Gly
                85
                                    90
Leu Ser Ser Gly Trp Leu Ala Asp Lys Val Gly Arg Arg Gly Ala Leu
                                105
                                                    110
Phe Tyr Asn Asn Leu Leu Ala Leu Ala Ala Ala Leu Met Gly Leu
                            120
                                                125
Ala Lys Ser Val Gly Ala Tyr Pro Met Val Ile Leu Gly Arg Leu Ile
```

```
150
                                        155
Thr Glu Ile Ser Pro Asn Asn Leu Arg Gly Met Leu Gly Ser Leu His
                                    170
                165
Gln Leu Leu Val Thr Ile Ala Ile Leu Val Ser Gln Ile Phe Gly Leu
                                185
                                                    190
Pro His Leu Leu Gly Thr Gly Asp Arg Trp Pro Leu Ile Phe Ala Phe
                            200
        195
                                                205
Thr Val Val Pro Ala Val Leu Gln Leu Ala Leu Leu Met Leu Cys Pro
                        215
                                            220
Glu Ser Pro Lys Tyr Thr Met Ala Val Arg Gly Gln Arg Asn Glu Ala
                    230
                                        235
Glu Ser Ala Leu Lys Lys Leu Arg Asp Thr Glu Asp Val Ser Thr Glu
                245
                                    250
Ile Glu Ala Met Gln Glu Glu Ala Thr Ala Ala Gly Val Gln Glu Lys
                                265
Pro Lys Met Gly Asp Met Phe Lys Gly Ala Leu Leu Trp Pro Met Ser
        275
                            280
                                                285
Ile Ala Ile Met Met Leu Ala Gln Gln Leu Ser Gly Ile Asn Val
                        295
                                             300
Ala Met Phe Tyr Ser Thr Val Ile Phe Arg Gly Ala Gly Leu Thr Gly
                    310
                                        315
Asn Glu Pro Phe Tyr Ala Thr Ile Gly Met Gly Ala Val Asn Val Ile
                325
                                    330
Met Thr Leu Ile Ser Val Trp Leu Val Asp His Pro Lys Phe Gly Arg
                                345
Arg Ser Leu Leu Leu Ala Gly Leu Thr Gly Met Phe Val Ser Thr Leu
        355
                            360
Leu Leu Val Gly Ala Leu Thr Ile Gln Asn Ser Gly Gly Asp Lys Trp
                        375
                                            380
Ala Ser Tyr Ser Ala Ile Gly Phe Val Leu Leu Phe Val Ile Ser Phe
                    390
                                        395
Ala Thr Gly Pro Gly Ala Ile Pro Trp Phe Phe Val Ser Glu Ile Phe
                405
                                    410
Asp Ser Ser Ala Arg Gly Asn Ala Asn Ser Ile Ala Val Met Val Asn
                                425
Trp Ala Ala Asn Leu Leu Val Gly Leu Thr Phe Leu Pro Ile Asn Asn
                            440
Leu Met Gln Gln Tyr Ser Phe Phe Ile Phe Ser Gly Phe Leu Ala Phe
                        455
                                             460
Phe Ile Phe Tyr Thr Trp Lys Phe Val Pro Glu Thr Lys Gly Lys Ser
                    470
                                        475
Ile Glu Gln Ile Gln Ala Glu Phe Glu Lys Arg Lys
                485
                                    490
<210> 209
<211> 22
<212> PRT
<213> Caenorhabditis elegans
```

Ile Gly Leu Asn Cys Gly Phe Ser Ser Ala Leu Val Pro Met Phe Leu

<400> 209

Arg Asn Glu Ala Glu Ser Ala Leu Lys Lys Leu Arg Asp Thr Glu Asp 1 5 10 15 Val Ser Thr Glu Ile Glu 20

<210> 210

```
<211> 28
<212> DNA
<213> Caenorhabditis elegans
<400> 210
tctcgttgtt tgccgtcgga tgtctgcc
<210> 211
<211> 223
<212> PRT
<213> Ascoris suum
<400> 211
Ala Lys Asn Asn Gly Glu Phe Val Arg Cys Val His Ser Val Gly Gln
                                    10
Pro Lys Pro Val Ala Thr Lys Val Ile Asn His Trp Pro Cys Asn Pro
            20
Glu Lys Thr Ile Ile Ala His Arg Pro Ala Glu Arg Glu Ile Trp Ser
        35
                            40
Phe Gly Ser Gly Tyr Gly Gly Asn Ser Leu Leu Gly Lys Lys Cys Phe
                        55
                                             60
Ala Leu Arg Ile Ala Met Asn Ile Gly Tyr Asp Glu Gly Trp Met Ala
                    70
                                         75
Glu His Met Leu Ile Met Gly Val Thr Ser Pro Lys Gly Glu Glu Arg
                85
                                    90
Phe Val Ala Ala Ala Phe Pro Ser Ala Cys Gly Lys Thr Asn Leu Ala
                                105
            100
                                                     110
Met Leu Glu Pro Thr Ile Pro Gly Trp Lys Val Arg Val Ile Gly Asp
        115
                            120
Asp Ile Ala Trp Met Lys Phe Gly Ala Asp Gly Arg Leu Tyr Ala Ile
                        135
                                             140
Asn Pro Glu Tyr Gly Phe Phe Gly Val Ala Pro Gly Thr Ser His Lys
                    150
                                         155
                                                             160
Thr Asn Pro Met Ala Met Ala Ser Phe Gln Glu Asn Thr Ile Phe Thr
                165
                                    170
                                                         175
Asn Val Ala Glu Thr Ala Asp Gly Glu Tyr Phe Trp Glu Gly Leu Glu
            180
                                185
                                                     190
His Glu Val Lys Asn Pro Lys Val Asp Met Ile Asn Trp Leu Gly Glu
                            200
                                                 205
Pro Trp His Ile Gly Asp Glu Ser Lys Ala Ala His Pro Asn Ser
    210
                        215
<210> 212
<211> 176
<212> PRT
<213> Caenorhabditis elegans or Ascoris suum
<400> 212
Ala Asn Phe Val Arg Cys His Ser Val Gly Pro Pro Val Val Ile Asn
1
                                    10
His Trp Pro Cys Asn Pro Glu Ile Ala His Arg Pro Glu Arg Glu Ile
            20
Trp Ser Phe Gly Ser Gly Tyr Gly Gly Asn Ser Leu Leu Gly Lys Lys
                            40
Cys Phe Ala Leu Arg Ile Ala Asn Ile Asp Glu Gly Trp Met Ala Glu
                        55
His Met Leu Ile Met Gly Val Thr Pro Gly Glu Phe Ala Ala Ala Phe
```

```
Pro Ser Ala Cys Gly Lys Thr Asn Leu Ala Met Leu Glu Pro Thr Pro
                85
                                    90
Gly Trp Lys Val Arg Gly Asp Asp Ile Ala Trp Met Lys Phe Gly Asp
                                105
            100
                                                     110
Gly Arg Leu Tyr Ala Ile Asn Pro Glu Gly Phe Phe Gly Val Ala Pro
                            120
                                                 125
Gly Thr Ser Lys Thr Asn Pro Met Ala Ala Phe Gln Asn Ile Phe Thr
                        135
Asn Val Ala Glu Thr Ala Gly Glu Tyr Phe Trp Glu Gly Leu Glu Glu
                    150
                                        155
Val Asp Trp Leu Gly Glu Trp His Ile Gly Ala Ala His Pro Asn Ser
                                    170
<210> 213
<211> 223
<212> PRT
<213> Caenorhabditis elegans
```

<400> 213 Ala Leu Gly Asn Gln Asp Phe Val Arg Cys Ile His Ser Val Gly Leu 10 Pro Arg Pro Val Lys Gln Arg Val Ile Asn His Trp Pro Cys Asn Pro 30 Glu Arg Val Leu Ile Ala His Arg Pro Pro Glu Arg Glu Ile Trp Ser Phe Gly Ser Gly Tyr Gly Gly Asn Ser Leu Leu Gly Lys Lys Cys Phe 55 Ala Leu Arg Ile Ala Ser Asn Ile Ala Lys Asp Glu Gly Trp Met Ala 70 75 Glu His Met Leu Ile Met Gly Val Thr Arg Pro Cys Gly Arg Glu His 90 Phe Ile Ala Ala Ala Phe Pro Ser Ala Cys Gly Lys Thr Asn Leu Ala 105 Met Leu Glu Pro Thr Leu Pro Gly Trp Lys Val Arg Cys Val Gly Asp 120 Asp Ile Ala Trp Met Lys Phe Gly Glu Asp Gly Arg Leu Tyr Ala Ile 135 Asn Pro Glu Ala Gly Phe Phe Gly Val Ala Pro Gly Thr Ser Asn Lys 150 155 Thr Asn Pro Met Ala Val Ala Thr Phe Gln Lys Asn Ser Ile Phe Thr 165 170 Asn Val Ala Glu Thr Ala Asn Gly Glu Tyr Phe Trp Glu Gly Leu Glu 185 190 Asp Glu Ile Ala Asp Lys Asn Val Asp Ile Thr Thr Trp Leu Gly Glu 200 Lys Trp His Ile Gly Glu Pro Gly Val Ala Ala His Pro Asn Ser

```
<210> 214
<211> 173
<212> PRT
<213> Ascoris suum
```

(21)/ ASCOLIS Sudi

<400> 214

Lys Gly Asp Phe Val Ser Leu Pro Lys His Val Gln Arg Phe Val Ala 1 5 10 15 Glu Lys Ala Glu Leu Met Lys Pro Ser Ala Ile Phe Ile Cys Asp Gly

```
25
Ser Gln Asn Glu Ala Asp Glu Leu Ile Ala Arg Cys Val Glu Arg Gly
                            40
Val Leu Val Pro Leu Lys Ala Tyr Lys Asn Asn Tyr Leu Cys Arg Thr
                        55
Asp Pro Arg Asp Val Ala Arg Val Glu Ser Lys Thr Trp Met Ile Thr
                                         75
Pro Glu Lys Tyr Asp Ser Val Cys His Thr Pro Glu Gly Val Lys Pro
                8.5
                                    90
Met Met Gly Gln Trp Met Ser Pro Asp Glu Phe Gly Lys Glu Leu Asp
            100
                                105
                                                    110
Asp Arg Phe Pro Gly Cys Met Ala Gly Arg Thr Met Tyr Val Ile Pro
        115
                            120
                                                125
Tyr Ser Met Gly Pro Val Gly Gly Pro Leu Ser Lys Ile Gly Ile Glu
    130
                        135
                                            140
Leu Thr Asp Ser Asp Tyr Val Val Leu Cys Met Arg Ile Met Thr Arg
                   150
                                        155
Met Gly Glu Pro Val Leu Lys Ala Leu Ala Lys Asn Asn
                165
<210> 215
<211> 120
<212> PRT
<213> Caenorhabditis elegans or Ascoris suum
<400> 215
Gly Asp Phe Leu Pro Val Gln Arg Phe Ala Glu Lys Ala Glu Leu Met
                                    10
Pro Ile Phe Ile Cys Asp Gly Ser Gln Glu Ala Asp Glu Leu Ile Glu
                                25
Arg Gly Leu Leu Ala Tyr Asn Asn Tyr Cys Arg Thr Asp Pro Asp Val
                            40
Ala Arg Val Glu Ser Lys Thr Trp Met Thr Lys Tyr Asp Val His Thr
                        55
Glu Gly Val Pro Met Gly Trp Pro Glu Leu Asp Arg Phe Pro Gly Cys
                    70
                                        7.5
Met Ala Gly Arg Met Tyr Val Ile Pro Ser Met Gly Pro Val Gly Gly
                                    90
Pro Leu Ser Lys Ile Gly Ile Leu Thr Asp Ser Tyr Val Val Leu Met
            100
                                105
Arg Ile Met Thr Arg Val Ala Leu
        115
<210> 216
<211> 173
<212> PRT
<213> Caenorhabditis elegans
<400> 216
Gln Gly Asp Phe His Leu Leu Pro Ala Lys Val Gln Arg Phe Ile Ala
Glu Lys Ala Glu Leu Met Arg Pro Arg Gly Ile Phe Ile Cys Asp Gly
                                25
Ser Gln His Glu Ala Asp Glu Leu Ile Asp Lys Leu Ile Glu Arg Gly
                            40
Met Leu Ser Lys Leu Glu Ala Tyr Glu Asn Asn Tyr Ile Cys Arg Thr
```

Asp Pro Lys Asp Val Ala Arg Val Glu Ser Lys Thr Trp Met Val Thr 75 Lys Asn Lys Tyr Asp Thr Val Thr His Thr Lys Glu Gly Val Glu Pro 85 90 Ile Met Gly His Trp Leu Ala Pro Glu Asp Leu Ala Thr Glu Leu Asp 100 105 110 Ser Arg Phe Pro Gly Cys Met Ala Gly Arg Ile Met Tyr Val Ile Pro 125 115 120 Phe Ser Met Gly Pro Val Gly Gly Pro Leu Ser Lys Ile Gly Ile Gln 135 140 Leu Thr Asp Ser Asn Tyr Val Val Leu Ser Met Arg Ile Met Thr Arg 150 155 160 Val Asn Asn Asp Val Trp Asp Ala Leu Gly Asn Gln Asp 165 170

<210> 217 <211> 107

<212> PRT

<213> Ascoris suum

<400> 217

Arg Phe Thr Ala Pro Ala Gly Gln Cys Pro Ile Ile His Pro Asp Trp Glu Lys Pro Glu Gly Val Pro Ile Asp Ala Ile Ile Phe Gly Gly Arg 25 30 Arg Pro Glu Gly Val Pro Leu Val Phe Glu Ser Arg Ser Trp Val His 35 Gly Ile Phe Val Gly Ala Cys Val Lys Ser Glu Ala Thr Ala Ala Ala 55 Glu His Thr Gly Lys Gln Val Met His Asp Pro Met Ala Met Arg Pro 70 75 Phe Met Gly Tyr Asn Phe Gly Arg Tyr Met Arg His Trp Met Lys Leu 90 95 Gly Gln Pro Pro His Lys Val Pro Lys Ile Phe 100 105

<210> 218 <211> 77

<212> PRT

<213> Caenorhabditis elegans or Ascoris suum

<400> 218

 Arg Phe Ala Pro Ala Gln Cys
 Pro Ile Ile His Pro Asp Trp Glu Pro 1
 5
 10
 15
 15

 Gly Val Pro Ile Ala Ile Ile Phe Gly Gly Arg Arg Pro 30
 20
 25
 30
 25
 30
 20

 Leu Glu Ser Trp His Gly Phe Gly Cys Lys Ser Glu Ala Thr Ala Ala 35
 40
 45
 45

 Ala Glu Thr Gly Lys Val Met His Asp Pro Met Ala Met Arg Pro Phe 50
 55
 60

 Met Gly Tyr Asn Phe Gly Tyr His Trp Leu Lys Val Phe 65
 70
 75

<210> 219

<211> 107

<212> PRT

<213> Caenorhabditis elegans

Ala Gln Arg Leu

```
<400> 219
Arg Phe Ala Ala Pro Ala Asn Gln Cys Pro Ile Ile His Pro Asp Trp
Glu Ser Pro Gln Gly Val Pro Ile Glu Ala Ile Ile Phe Gly Gly Arg
Arg Pro Gln Gly Val Pro Leu Ile Tyr Glu Thr Asn Ser Trp Glu His
Gly Val Phe Thr Gly Ser Cys Leu Lys Ser Glu Ala Thr Ala Ala Ala
                        55
                                            60
Glu Phe Thr Gly Lys Thr Val Met His Asp Pro Met Ala Met Arg Pro
                    70
                                        7.5
Phe Met Gly Tyr Asn Phe Gly Lys Tyr Leu Gln His Trp Leu Asp Leu
                                    90
                85
Lys Thr Asp Ser Arg Lys Val Ile Asp Phe Phe
            100
<210> 220
<211> 116
<212> PRT
<213> Ascoris suum
<400> 220
Val Pro Lys Ile Phe His Val Asn Trp Phe Arg Gln Ser Ala Asp His
Lys Phe Leu Trp Pro Gly Tyr Gly Asp Asn Ile Arg Val Ile Asp Trp
            20
                                25
Ile Leu Arg Arg Cys Ser Gly Asp Ala Thr Ile Ala Glu Glu Thr Pro
                            40
Ile Gly Phe Ile Pro Lys Lys Gly Thr Ile Asn Leu Glu Gly Leu Pro
                        55
Asn Val Asn Trp Asp Glu Leu Met Ser Ile Pro Lys Ser Tyr Trp Leu
                    70
                                        75
Glu Asp Met Val Glu Thr Lys Thr Phe Phe Glu Asn Gln Val Gly Ser
                                    90
                85
Asp Leu Pro Pro Glu Ile Ala Lys Glu Leu Glu Ala Gln Thr Glu Arg
            100
                                105
Ile Lys Ala Leu
        115
<210> 221
<211> 68
<212> PRT
<213> Caenorhabditis elegans or Ascoris suum
<400> 221
Pro Lys Ile His Val Asn Trp Phe Arg Lys Phe Leu Trp Pro Gly Gly
1
Asp Asn Ile Arg Val Ile Asp Trp Ile Arg Arg Gly Ile Glu Thr Pro
            20
Ile Gly Pro Lys Gly Ile Asn Leu Glu Gly Leu Val Asn Trp Asp Glu
                            40
Leu Met Ser Pro Tyr Trp Asp Glu Phe Gln Val Gly Asp Leu Pro Glu
                        55
   50
```

```
<210> 222
<211> 116
<212> PRT
<213> Caenorhabditis elegans
<400> 222
Met Pro Lys Ile Tyr His Val Asn Trp Phe Arg Lys Asp Ser Asn Asn
                                    10
Lys Phe Leu Trp Pro Gly Phe Gly Asp Asn Ile Arg Val Ile Asp Trp
                                25
Ile Ile Arg Arg Leu Asp Gly Glu Gln Glu Ile Gly Val Glu Thr Pro
        35
Ile Gly Thr Val Pro Ala Lys Gly Ser Ile Asn Leu Glu Gly Leu Gly
                        55
Glu Val Asn Trp Asp Glu Leu Met Ser Val Pro Ala Asp Tyr Trp Lys
                    70
                                        75
Gln Asp Ala Gln Glu Ile Arg Lys Phe Leu Asp Glu Gln Val Gly Glu
                85
                                    90
Asp Leu Pro Glu Pro Val Arg Ala Glu Met Asp Ala Gln Glu Lys Arg
            100
                                105
Val Gln Thr Leu
        115
<210> 223
<211> 36
<212> PRT
<213> Ascoris suum
<400> 223
Ser Leu Ser His Phe Lys Asp Asp Phe Ala Val Val Ser Glu Val
                                    1.0
Val Thr His Lys Gln Asn His Ile Pro Val Ile Lys Gly Asp Phe Val
            20
                                25
Ser Leu Pro Lys
        35
<210> 224
<211> 15
<212> PRT
<213> Caenorhabditis elegans or Ascoris suum
<400> 224
Ser Leu Asp Phe Val Val Glu Val Val His Pro Lys Phe Ser Lys
                                    10
<210> 225
<211> 36
<212> PRT
<213> Caenorhabditis elegans
<400> 225
Ser Leu Arg Gln Ile Ser Glu Asp Ala Phe Tyr Val Val Asn Glu Val
Val Met Lys Arg Leu Gly His Val Pro Ile Leu Lys Val Ile Phe Glu
```

```
Ser Ser Glu Lys
        35
<210> 226
<211> 25
<212> PRT
<213> Ascoris suum
<400> 226
Gly Cys Met Ala Gly Arg Thr Met Tyr Val Ile Pro Tyr Ser Met Gly
                5
Pro Val Gly Gly Pro Leu Ser Lys Ile
            20
<210> 227
<211> 9
<212> PRT
<213> Caenorhabditis elegans or Ascoris suum
<400> 227
Gly Cys Arg Val Pro Ser Pro Leu Lys
                5
<210> 228
<211> 25
<212> PRT
<213> Caenorhabditis elegans
<400> 228
Gly Cys Ser Gly Arg Arg Val Leu Cys Val Cys Pro Cys Ser His Ser
                5
Ser Ser Ala Leu Pro Leu Gln Lys Val
            20
<210> 229
<211> 16
<212> PRT
<213> Ascoris suum
<400> 229
Leu Pro Asn Val Asn Trp Asp Glu Leu Met Ser Ile Pro Lys Ser Tyr
                5
                                    10
<210> 230
<211> 7
<212> PRT
<213> Caenorhabditis elegans or Ascoris suum
<400> 230
Leu Asn Trp Ser Pro Ser Tyr
                 5
<210> 231
```

```
<211> 16
<212> PRT
<213> Caenorhabditis elegans
<400> 231
Leu Glu Ser Phe Asn Trp Phe Ser Phe Val Ser Cys Pro Asp Ser Tyr
                                     10
<210> 232
<211> 14
<212> PRT
<213> Ascoris suum
<400> 232
Ser Val Cys His Thr Pro Glu Gly Val Lys Pro Met Met Gly
                                     10
<210> 233
<211> 6
<212> PRT
<213> Caenorhabditis elegans or Ascoris suum
<400> 233
Val His Pro Pro Met Gly
1
<210> 234
<211> 14
<212> PRT
<213> Caenorhabditis elegans
<400> 234
Thr Val Met His Asp Pro Met Ala Met Arg Pro Phe Met Gly
1
                                     10
<210> 235
<211> 197
<212> PRT
<213> Homo sapiens
<400> 235
Ser Gly Phe Phe Asp Tyr Gly Ser Phe Ser Glu Ile Met Gln Pro Trp
1
Ala Gln Thr Val Val Val Gly Arg Ala Arg Leu Gly Gly Ile Pro Val
                                                     30
            20
                                 25
Gly Val Val Ala Val Glu Thr Arg Thr Val Glu Leu Ser Val Pro Ala
        35
                            40
Asp Pro Ala Asn Leu Asp Ser Glu Ala Lys Ile Ile Gln Gln Ala Gly
                        55
    50
                                             60
Gln Val Trp Phe Pro Asp Ser Ala Phe Lys Thr Tyr Gln Ala Ile Lys
                    70
                                         75
Asp Phe Asn Arg Glu Gly Leu Pro Leu Met Val Phe Ala Asn Trp Arg
                                     90
Gly Phe Ser Gly Gly Met Lys Asp Met Tyr Asp Gln Val Leu Lys Phe
            100
                                 105
                                                     110
```

```
Gly Ala Tyr Ile Val Asp Gly Leu Arg Glu Cys Ser Gln Pro Val Met
        115
                            120
Val Tyr Ile Pro Pro Gln Ala Glu Leu Arg Gly Gly Ser Trp Val Val
                        135
                                             140
Ile Asp Pro Thr Ile Asn Pro Arg His Met Glu Met Tyr Ala Asp Arg
                    150
                                         155
Glu Ser Arg Gly Ser Val Leu Glu Pro Glu Gly Thr Val Glu Ile Lys
                                    170
                165
Phe Arg Lys Lys Asp Leu Val Lys Thr Met Arg Arg Val Asp Pro Val
                                185
            180
Tyr Ile Arg Leu Ala
        195
<210> 236
<211> 109
<212> PRT
<213> Caenorhabditis elegans or Homo sapiens
<400> 236
Gly Asp Ser Phe Glu Ile Trp Ala Val Gly Arg Ala Arg Leu Gly Ile
                                    1.0
Pro Gly Val Val Glu Arg Val Pro Ala Asp Pro Ala Ser Gln Ala Gly
                                25
Gln Val Trp Pro Asp Ser Ala Phe Lys Thr Ala Ile Asp Asn Glu Leu
        35
                            40
                                                 4.5
Pro Leu Met Ala Arg Gly Phe Ser Gly Gly Lys Asp Met Tyr Asp Val
                        55
                                             60
Leu Lys Phe Gly Ala Ile Val Asp Leu Pro Val Val Tyr Ile Pro Glu
                    70
                                        75
Leu Arg Gly Gly Trp Val Asp Ile Pro Ala Asp Ser Arg Gly Leu Glu
                8.5
                                    90
Pro Val Ile Lys Phe Arg Lys Met Arg Asp Pro Tyr Leu
                                105
<210> 237
<211> 197
<212> PRT
<213> Caenorhabditis elegans
<400> 237
Thr Gly Ile Cys Asp Thr Met Ser Phe Asp Glu Ile Cys Gly Asp Trp
                                    10
Ala Lys Ser Ile Val Ala Gly Arg Ala Arg Leu Cys Gly Ile Pro Ile
                                25
Gly Val Val Ser Ser Glu Phe Arg Asn Phe Ser Thr Ile Val Pro Ala
                            40
Asp Pro Ala Ile Asp Gly Ser Gln Val Gln Asn Thr Gln Arg Ala Gly
                        55
                                             60
Gln Val Trp Tyr Pro Asp Ser Ala Phe Lys Thr Ala Glu Ala Ile Asn
                    70
                                        75
Asp Leu Asn Lys Glu Asn Leu Pro Leu Met Ile Ile Ala Ser Leu Arg
                85
                                    90
Gly Phe Ser Gly Gly Gln Lys Asp Met Tyr Asp Met Val Leu Lys Phe
```

105 Gly Ala Gln Ile Val Asp Ala Leu Ala Val Tyr Asn Arg Pro Val Ile 120

Val Tyr Ile Pro Glu Ala Gly Glu Leu Arg Gly Gly Ala Trp Ala Val

```
130
                       135
Leu Asp Ser Lys Ile Arg Pro Glu Phe Ile His Leu Val Ala Asp Glu
                   150
                                      155
Lys Ser Arg Gly Gly Ile Leu Glu Pro Asn Ala Val Val Gly Ile Lys
               165
                                  170
                                                      175
Phe Arg Lys Pro Met Met Met Glu Met Met Lys Arg Ser Asp Pro Thr
           180
                              185
                                                  190
Tyr Ser Lys Leu Ser
       195
<210> 238
<211> 124
<212> PRT
<213> Homo sapiens
<220>
<221> VARIANT
<222> (1)...(124)
<223> Xaa = Any Amino Acid
<400> 238
Val Gly Tyr Pro Val Met Ile Lys Ala Ser Glu Gly Gly Gly Lys
Gly Ile Arg Lys Val Asn Asn Ala Asp Asp Phe Pro Asn Leu Phe Arg
           20
                               25
Gln Val Gln Ala Glu Val Pro Gly Ser Pro Ile Phe Val Met Arg Leu
                           40
Ala Lys Gln Ser Arg His Leu Glu Val Gln Ile Leu Ala Asp Gln Tyr
                       55
Gly Asn Ala Ile Ser Leu Phe Gly Arg Asp Cys Ser Val Gln Arg Arg
                   70
                                      75
90
Val Phe Glu His Met Glu Gln Cys Ala Val Lys Leu Ala Lys Met Val
                              105
Gly Tyr Val Ser Ala Gly Thr Val Glu Tyr Leu Tyr
       115
<210> 239
<211> 68
<212> PRT
<213> Homo sapiens or Caenorhabditis elegans
<400> 239
Gly Pro Met Ile Lys Ala Ser Glu Gly Gly Gly Lys Gly Ile Arg
1
                                  10
Lys Asp Phe Phe Val Glu Val Gly Ser Pro Ile Phe Met Arg His Glu
                               25
Val Gln Leu Ala Asp Tyr Asn Ile Ser Arg Asp Cys Ser Gln Arg Arg
                           40
Gln Lys Met Ala Val Leu Ala Lys Val Gly Tyr Ser Ala Gly Thr Val
Glu Tyr Leu Tyr
```

```
<211> 124
<212> PRT
<213> Caenorhabditis elegans
<400> 240
Ile Gly Phe Pro Leu Met Ile Lys Ala Ser Glu Gly Gly Gly Lys
Gly Ile Arg Lys Cys Thr Lys Val Glu Asp Phe Lys Ser Met Phe Glu
            20
                                25
Glu Val Ala Gln Glu Val Gln Gly Ser Pro Ile Phe Leu Met Lys Cys
                            40
Val Asp Gly Ala Arg His Ile Glu Val Gln Leu Leu Ala Asp Arg Tyr
                        55
Glu Asn Val Ile Ser Val Tyr Thr Arg Asp Cys Ser Ile Gln Arg Arg
                    70
                                        75
Cys Gln Lys Ile Ile Glu Glu Ala Pro Ala Ile Ile Ala Ser Ser His
                                    90
                8.5
Ile Arg Lys Ser Met Gln Glu Asp Ala Val Arg Leu Ala Lys Tyr Val
                                105
Gly Tyr Glu Ser Ala Gly Thr Val Glu Tyr Leu Tyr
                            120
<210> 241
<211> 116
<212> PRT
<213> Rat
<400> 241
Lys Glu Glu Gly Leu Gly Ala Glu Asn Leu Arg Gly Ser Gly Met Ile
                                    1.0
Ala Gly Glu Ser Ser Leu Ala Tyr Asp Glu Ile Ile Thr Ile Ser Leu
                                25
Val Thr Cys Arg Ala Ile Gly Ile Gly Ala Tyr Leu Val Arg Leu Gly
       35
                            40
Gln Arg Thr Ile Gln Val Glu Asn Ser His Leu Ile Leu Thr Gly Ala
                        55
Gly Ala Leu Asn Lys Val Leu Gly Arg Glu Val Tyr Thr Ser Asn Asn
                    70
                                        7.5
Gln Leu Gly Gly Ile Gln Ile Met His Asn Asn Gly Val Thr His Cys
                85
                                    90
Thr Val Cys Asp Asp Phe Glu Gly Val Phe Thr Val Leu His Trp Leu
            100
                                105
Ser Tyr Met Pro
       115
<210> 242
<211> 65
<212> PRT
<213> Caenorhabditis elegans or Rat
<400> 242
Lys Glu Gly Glu Asn Leu Gly Ser Gly Ile Ala Gly Glu Ala Tyr Glu
Thr Val Thr Arg Gly Ile Gly Ala Tyr Arg Leu Arg Gln Ser His Leu
            20
Ile Leu Thr Gly Ala Leu Asn Leu Gly Val Tyr Thr Ser Asn Asn Gln
```

```
Leu Gly Gly Met Asn Gly Val Thr His Val Asp Glu Gly Val Trp Ser
    50
                        55
Pro
65
<210> 243
<211> 116
<212> PRT
<213> Caenorhabditis elegans
<400> 243
Lys Asn Glu Lys Ile Gly Val Glu Asn Leu Gln Gly Ser Gly Leu Ile
1
Ala Gly Glu Thr Ala Arg Ala Tyr Ala Glu Val Pro Thr Tyr Cys Tyr
            20
                                25
Val Thr Gly Arg Ser Val Gly Ile Gly Ala Tyr Thr Ala Arg Leu Ala
        35
His Arg Ile Val Gln His Lys Gln Ser His Leu Ile Leu Thr Gly Tyr
    50
                        55
Glu Ala Leu Asn Thr Leu Leu Gly Lys Lys Val Tyr Thr Ser Asn Asn
                    70
Gln Leu Gly Gly Pro Glu Val Met Phe Arg Asn Gly Val Thr His Ala
                85
                                     90
Val Val Asp Asn Asp Leu Glu Gly Ile Ala Lys Val Ile Arg Trp Met
            100
                                105
                                                     110
Ser Phe Leu Pro
        115
<210> 244
<211> 119
<212> PRT
<213> Homo sapiens
<400> 244
His Val Ile Ala Ala Arg Ile Thr Ser Glu Asn Pro Asp Glu Gly Phe
1
                                     10
Lys Pro Ser Ser Gly Thr Val Gln Glu Leu Asn Phe Arg Ser Asn Lys
Asn Val Trp Gly Tyr Phe Ser Val Ala Ala Ala Gly Gly Leu His Glu
                            40
Phe Ala Asp Ser Gln Phe Gly His Cys Phe Ser Trp Gly Glu Asn Arg
                        55
Glu Glu Ala Ile Ser Asn Met Val Val Ala Leu Lys Glu Leu Ser Ile
                    70
                                         75
Arg Gly Asp Phe Arg Thr Thr Val Glu Tyr Leu Ile Lys Leu Leu Glu
                85
                                     90
Thr Glu Ser Phe Gln Leu Asn Arg Ile Asp Thr Gly Trp Leu Asp Arg
            100
                                105
Leu Ile Ala Glu Lys Val Gln
        115
<210> 245
<211> 59
<212> PRT
<213> Caenorhabditis elegans or Homo sapiens
```

```
<400> 245
His Ile Ala Ala Arg Ile Thr Glu Asn Pro Asp Phe Pro Ser Gly Val
Glu Asn Phe Ser Trp Tyr Phe Ser Val His Phe Ala Asp Ser Gln Phe
Gly His Phe Gly Arg Glu Ala Met Leu Lys Ile Arg Phe Thr Val Tyr
Leu Leu Phe Asn Thr Trp Leu Asp Ile Ala Lys
<210> 246
<211> 119
<212> PRT
<213> Caenorhabditis elegans
<400> 246
His Ala Ile Ala Arg Ile Thr Cys Glu Asn Pro Asp Asp Ser Phe
1
Arg Pro Ser Thr Gly Lys Val Tyr Glu Ile Asn Phe Pro Ser Ser Gln
                                25
Asp Ala Trp Ala Tyr Phe Ser Val Gly Arg Gly Ser Ser Val His Gln
                            40
Phe Ala Asp Ser Gln Phe Gly His Ile Phe Thr Arg Gly Thr Ser Arg
                        55
Thr Glu Ala Met Asn Thr Met Cys Ser Thr Leu Lys His Met Thr Ile
                    70
                                        75
Arg Ser Ser Phe Pro Thr Gln Val Asn Tyr Leu Val Asp Leu Met His
                                    90
Asp Ala Asp Phe Ile Asn Asn Ala Phe Asn Thr Gln Trp Leu Asp Lys
            100
                                105
Arg Ile Ala Met Lys Ile Lys
       115
<210> 247
<211> 90
<212> PRT
<213> Rat
<400> 247
Pro Gly Gly Ala Asn Asn Asn Tyr Ala Asn Val Glu Leu Ile Leu
Asp Ile Ala Lys Arg Ile Pro Val Gln Ala Val Trp Ala Gly Trp Gly
                                25
His Ala Ser Glu Asn Pro Lys Leu Pro Glu Leu Leu Leu Lys Asn Gly
Ile Ala Phe Met Gly Pro Pro Ser Gln Ala Met Trp Ala Leu Gly Asp
                        55
Lys Ile Ala Ser Ser Ile Val Ala Gln Thr Ala Gly Ile Pro Thr Leu
Pro Trp Ser Gly Ser Gly Leu Arg Val Asp
<210> 248
<211> 55
<212> PRT
```

<213> Caenorhabditis elegans or Rat

```
<400> 248
Pro Gly Asn Asn Asn Ala Asn Val Ile Leu Ala Val Ala Val Trp Ala
Gly Trp Gly His Ala Ser Glu Asn Pro Leu Pro Leu Ile Ala Phe Gly
            20
Pro Pro Ala Met Leu Gly Asp Lys Ile Ala Ser Ile Ala Gln Thr Gly
Pro Thr Trp Ser Gly Ser Gly
    50
<210> 249
<211> 90
<212> PRT
<213> Caenorhabditis elegans
<400> 249
Pro Ser Gly Thr Asn Lys Asn Asn Phe Ala Asn Val Asp Glu Ile Leu
Lys His Ala Ile Lys Tyr Glu Val Asp Ala Val Trp Ala Gly Trp Gly
His Ala Ser Glu Asn Pro Asp Leu Pro Arg Arg Leu Asn Asp His Asn
Ile Ala Phe Ile Gly Pro Pro Ala Ser Ala Met Phe Ser Leu Gly Asp
Lys Ile Ala Ser Thr Ile Ile Ala Gln Thr Val Gly Val Pro Thr Val
                    70
Ala Trp Ser Gly Ser Gly Ile Thr Met Glu
                85
<210> 250
<211> 67
<212> PRT
<213> Caenorhabditis elegans
<400> 250
Val Ile Lys Asn Leu Gly Tyr Met Val Asp Asn His Gly Phe Val Pro
1
Asn Gly Gly Arg Val Tyr Tyr Leu Thr Arg Ser Gln Pro Pro Leu Leu
            20
Thr Pro Met Val Tyr Glu Tyr Tyr Met Ser Thr Gly Asp Leu Asp Phe
                            40
Val Met Glu Ile Leu Pro Thr Leu Asp Lys Glu Tyr Glu Phe Trp Ile .
                        55
                                             60
Lys Asn Arg
65
<210> 251
<211> 36
<212> PRT
<213> Caenorhabditis elegans
<400> 251
Ile Asn Gly Phe Val Pro Asn Gly Gly Arg Val Tyr Tyr Leu Arg Ser
                                    10
Gln Pro Pro Pro Met Val Tyr Glu Tyr Tyr Thr Asp Val Pro Lys Glu
```

```
Tyr Phe Trp Arg
        35
<210> 252
<211> 67
<212> PRT
<213> Caenorhabditis elegans
<400> 252
Met Ile Leu Asn Phe Ala His Ile Ile Glu Thr Tyr Gly Phe Val Pro
                                     10
                                                         15
Asn Gly Gly Arg Val Tyr Tyr Leu Arg Arg Ser Gln Pro Pro Phe Phe
                                25
Ala Pro Met Val Tyr Glu Tyr Tyr Leu Ala Thr Gln Asp Ile Gln Leu
                            40
                                                 45
Val Ala Asp Leu Ile Pro Val Ile Glu Lys Glu Tyr Thr Phe Trp Ser
Glu Arg Arg
65
<210> 253
<211> 92
<212> PRT
<213> Caenorhabditis elegans
<400> 253
Met Asp Ser Ile Arg Thr Trp Ser Ile Ile Pro Ala Asp Leu Asn Ala
                                     10
Phe Met Cys Ala Asn Ala Arg Ile Leu Ala Ser Leu Tyr Glu Ile Ala
                                25
                                                     30
Gly Asp Phe Lys Lys Val Lys Val Phe Glu Gln Arg Tyr Thr Trp Ala
                            40
                                                 45
Lys Arg Glu Met Arg Glu Leu His Trp Asn Glu Thr Asp Gly Ile Trp
                        55
Tyr Asp Tyr Asp Ile Glu Leu Lys Thr His Ser Asn Gln Tyr Tyr Val
                    70
                                         75
Ser Asn Ala Val Pro Leu Tyr Ala Lys Cys Tyr Asp
<210> 254
<211> 32
<212> PRT
<213> Caenorhabditis elegans
<400> 254
Ile Thr Ile Pro Asp Leu Asn Ala Phe Cys Asn Ile Tyr Gly Lys Arg
                                                         15
                                    10
Thr Trp Tyr Asp Tyr Thr His Ser Asn Ala Val Pro Leu Cys Tyr Asp
<210> 255
<211> 92
<212> PRT
<213> Caenorhabditis elegans
```

```
<400> 255
Ile Ser Thr Ile Glu Thr Thr Asn Ile Val Pro Val Asp Leu Asn Ala
                                     10
Phe Leu Cys Tyr Asn Met Asn Ile Met Gln Leu Phe Tyr Lys Leu Thr
            20
                                 25
                                                     30
Gly Asn Pro Leu Lys His Leu Glu Trp Ser Ser Arg Phe Thr Asn Phe
                            40
Arg Glu Ala Phe Thr Lys Val Phe Tyr Val Pro Ala Arg Lys Gly Trp
                        55
Tyr Asp Tyr Asn Leu Arg Thr Leu Thr His Asn Thr Asp Phe Phe Ala
                    70
Ser Asn Ala Val Pro Leu Phe Ser Gln Cys Tyr Asp
                85
<210> 256
<211> 102
<212> PRT
<213> Caenorhabditis elegans
<400> 256
Val His Asp Tyr Leu Glu Arg Gln Gly Leu Leu Lys Tyr Thr Lys Gly
1
                                     10
Leu Pro Thr Ser Leu Ala Met Ser Ser Thr Gln Gln Trp Asp Lys Glu
            20
                                 25
Asn Ala Trp Pro Pro Met Ile His Met Val Ile Glu Gly Phe Arg Thr
        35
                            40
                                                 45
Thr Gly Asp Ile Lys Leu Met Lys Val Ala Glu Lys Met Ala Thr Ser
                        55
                                             60
Trp Leu Thr Gly Thr Tyr Gln Ser Phe Ile Arg Thr His Ala Met Phe
                    70
                                         75
Glu Lys Tyr Asn Val Thr Pro His Thr Glu Glu Thr Ser Gly Gly Gly
                85
                                     90
Gly Gly Glu Tyr Glu Val
            100
<210> 257
<211> 37
<212> PRT
<213> Caenorhabditis elegans
<400> 257
Val Gly Gly Pro Thr Ser Gln Gln Trp Asp Asn Trp Pro Met His Met
                                    10
Ile Glu Gly Arg Leu Ala Ala Trp Leu Gln Phe Met Glu Lys Tyr Asn
            20
                                25
                                                     30
Val Gly Gly Glu Val
        35
<210> 258
<211> 102
<212> PRT
<213> Caenorhabditis elegans
<400> 258
Val Tyr Asn Glu Met Gln Asn Ser Gly Ala Phe Ser Ile Pro Gly Gly
                 5
                                     10
                                                         1.5
```

```
Ile Pro Thr Ser Met Asn Glu Glu Thr Asn Gln Gln Trp Asp Phe Pro
                                25
Asn Gly Trp Ser Pro Met Asn His Met Ile Ile Glu Gly Leu Arg Lys
                            40
Ser Asn Asn Pro Ile Leu Gln Gln Lys Ala Phe Thr Leu Ala Glu Lys
                        55
Trp Leu Glu Thr Asn Met Gln Thr Phe Asn Val Ser Asp Glu Met Trp
                    70
                                        75
Glu Lys Tyr Asn Val Lys Glu Pro Leu Gly Lys Leu Ala Thr Gly Gly
                85
Glu Tyr Glu Val Gln Val
            100
<210> 259
<211> 58
<212> PRT
<213> Caenorhabditis elegans
<400> 259
Tyr Gln Tyr Lys Ala Lys Leu Lys Val Pro Arg Pro Glu Ser Tyr Arg
                                     10
Glu Asp Ser Glu Leu Ala Glu His Leu Gln Thr Glu Ala Glu Lys Ile
                                25
            20
Gln Met Trp Ser Glu Ile Ala Ser Ala Ala Glu Thr Gly Trp Asp Phe
        35
Ser Thr Arg Trp Phe Ser Gln Asn Gly Asp
    50
<210> 260
<211> 29
<212> PRT
<213> Caenorhabditis elegans
<400> 260
Gln Tyr Pro Arg Pro Glu Ser Arg Glu Asp Ala Glu His Thr Lys Gln
                                    10
                                                         15
Ser Ala Ala Glu Gly Trp Asp Phe Ser Arg Trp Phe Asp
            20
<210> 261
<211> 58
<212> PRT
<213> Caenorhabditis elegans
<400> 261
Phe Gln Tyr Arg Thr Glu Ala Glu Thr Pro Arg Pro Glu Ser Phe Arg
                                     10
Glu Asp Val Leu Ser Ala Glu His Phe Thr Thr Lys Asp Arg Lys Lys
            20
                                25
Gln Phe Phe Lys Asp Leu Gly Ser Ala Ala Glu Ser Gly Trp Asp Phe
```

40

Ser Ser Arg Trp Phe Lys Asn His Lys Asp

<210> 262

```
<211> 21
<212> PRT
<213> Caenorhabditis elegans
<400> 262
Gln Thr Gly Phe Gly Trp Thr Asn Gly Val Ile Leu Asp Leu Leu Asp
1
Lys Tyr Gly Asp Gln
            20
<210> 263
<211> 13
<212> PRT
<213> Caenorhabditis elegans
<400> 263
Gln Gly Phe Gly Trp Thr Asn Gly Leu Asp Leu Tyr Asp
<210> 264
<211> 21
<212> PRT
<213> Caenorhabditis elegans
<400> 264
Gln Ala Gly Phe Gly Trp Thr Asn Gly Ala Ala Leu Asp Leu Ile Phe
1
                                     10
Thr Tyr Ser Asp Arg
            20
<210> 265
<211> 24
<212> PRT
<213> Caenorhabditis elegans
<400> 265
Ser Ser Ser Thr Ala Ser Lys Phe Ser Phe Ser Leu Ser Asn Ile Thr
1
                                     10
Phe Val Val Phe Ile Leu Tyr Ile
            20
<210> 266
<211> 10
<212> PRT
<213> Caenorhabditis elegans
<400> 266
Ser Ser Ser Phe Ser Val Phe Leu Tyr Ile
                 5
1
                                     10
<210> 267
<211> 24
<212> PRT
<213> Caenorhabditis elegans
```

```
<400> 267
Thr Ser Ser Ser Ser Thr Phe Gly Tyr Ser Asn Ile Leu Thr Leu
                5
                                    10
Ile Thr Val Phe Val Leu Tyr Ile
            20
<210> 268
<211> 7
<212> PRT
<213> Caenorhabditis elegans
<400> 268
Gly Gly Glu Tyr Glu Val Gln
<210> 269
<211> 7
<212> PRT
<213> Caenorhabditis elegans
<400> 269
Gly Gly Glu Tyr Glu Val Gln
<210> 270
<211> 7
<212> PRT
<213> Caenorhabditis elegans
<400> 270
Gly Gly Glu Tyr Glu Val Gln
                 5
<210> 271
<211> 18
<212> PRT
<213> Caenorhabditis elegans
<400> 271
Lys Thr His Ser Asn Gln Tyr Tyr Val Ser Asn Ala Val Pro Leu Tyr
1
                 5
                                    10
Ala Lys
<210> 272
<211> 8
<212> PRT
<213> Caenorhabditis elegans
<400> 272
Lys Tyr Tyr Val Ser Pro Tyr Lys
                 5
```

```
<210> 273
<211> 18
<212> PRT
<213> Caenorhabditis elegans
<400> 273
Lys Phe Thr Ala His Pro Tyr Tyr Val Ser Arg Thr Pro Pro Arg Tyr
1
                 5
His Lys
<210> 274
<211> 67
<212> PRT
<213> Caenorhabditis elegans
<400> 274
Val Ile Lys Asn Leu Gly Tyr Met Val Asp Asn His Gly Phe Val Pro
                                     10
Asn Gly Gly Arg Val Tyr Tyr Leu Thr Arg Ser Gln Pro Pro Leu Leu
Thr Pro Met Val Tyr Glu Tyr Tyr Met Ser Thr Gly Asp Leu Asp Phe
                            40
                                                 45
Val Met Glu Ile Leu Pro Thr Leu Asp Lys Glu Tyr Glu Phe Trp Ile
                        55
                                             60
Lys Asn Arg
65
<210> 275
<211> 43
<212> PRT
<213> Caenorhabditis elegans
<400> 275
Ile Asn Leu Met Val Asp Gly Phe Val Pro Asn Gly Gly Arg Val Tyr
                                     10
Tyr Leu Arg Ser Gln Pro Pro Leu Met Val Tyr Glu Tyr Thr Asp Phe
            20
Val Glu Leu Pro Thr Leu Lys Glu Phe Trp Arg
        35
                            40
<210> 276
<211> 67
<212> PRT
<213> Caenorhabditis elegans
<400> 276
Met Ile Arq Asn Leu Alâ Ser Met Val Asp Lys Tyr Gly Phe Val Pro
                                                         15
Asn Gly Gly Arg Val Tyr Tyr Leu Gln Arg Ser Gln Pro Pro Phe Leu
            20
Ala Ala Met Val Tyr Glu Leu Tyr Glu Ala Thr Asn Asp Lys Ala Phe
                            40
Val Ala Glu Leu Leu Pro Thr Leu Leu Lys Glu Leu Asn Phe Trp Asn
                                             60
```

Glu Lys Arg

```
<210> 277
<211> 84
<212> PRT
<213> Caenorhabditis elegans
<400> 277
Ile Ile Pro Ala Asp Leu Asn Ala Phe Met Cys Ala Asn Ala Arg Ile
                                     10
Leu Ala Ser Leu Tyr Glu Ile Ala Gly Asp Phe Lys Lys Val Lys Val
                                25
Phe Glu Gln Arg Tyr Thr Trp Ala Lys Arg Glu Met Arg Glu Leu His
                            40
Trp Asn Glu Thr Asp Gly Ile Trp Tyr Asp Tyr Asp Ile Glu Leu Lys
                        55
                                             60
Thr His Ser Asn Gln Tyr Tyr Val Ser Asn Ala Val Pro Leu Tyr Ala
                    70
Lys Cys Tyr Asp
<210> 278
<211> 31
<212> PRT
<213> Caenorhabditis elegans
<400> 278
Pro Asp Leu Asn Cys Asn Ile Leu Tyr Glu Gly Asp Lys Phe Asn Thr
                                    10
Asp Gly Trp Tyr Asp Tyr His Tyr Ser Ala Val Pro Leu Cys Tyr
<210> 279
<211> 84
<212> PRT
<213> Caenorhabditis elegans
<400> 279
Val Leu Pro Val Asp Leu Asn Gly Leu Leu Cys Trp Asn Met Asp Ile
                                     10
Met Glu Tyr Leu Tyr Glu Gln Ile Gly Asp Thr Lys Asn Ser Gln Ile
                                25
Phe Arg Asn Lys Arg Ala Asp Phe Arg Asp Thr Val Gln Asn Val Phe
                            40
Tyr Asn Arg Thr Asp Gly Thr Trp Tyr Asp Tyr Asn Leu Arg Thr Gln
                        55
                                             60
Ser His Asn Pro Arg Phe Tyr Thr Ser Thr Ala Val Pro Leu Phe Thr
                    70
Asn Cys Tyr Asn
<210> 280 ,
<211> 48
<212> PRT
```

<213> Caenorhabditis elegans

```
<400> 280
Tyr Leu Glu Arg Gln Gly Leu Leu Lys Tyr Thr Lys Gly Leu Pro Thr
                                     10
Ser Leu Ala Met Ser Ser Thr Gln Gln Trp Asp Lys Glu Asn Ala Trp
                                25
Pro Pro Met Ile His Met Val Ile Glu Gly Phe Arg Thr Thr Gly Asp
                            40
<210> 281
<211> 20
<212> PRT
<213> Caenorhabditis elegans
<400> 281
Gly Tyr Gly Pro Thr Ser Ser Gln Gln Trp Asp Asn Trp Pro His Met
1
                                    10
Ile Glu Gly Arg
            20
<210> 282
<211> 48
<212> PRT
<213> Caenorhabditis elegans
<400> 282
Phe Phe Gln Lys Met Gly Val Phe Thr Tyr Pro Gly Gly Ile Pro Thr
1
                                     10
Ser Met Ser Gln Glu Ser Asp Gln Gln Trp Asp Phe Pro Asn Gly Trp
            20
                                25
Ser Pro Asn Asn His Met Ile Ile Glu Gly Leu Arg Lys Ser Ala Asn
                            40
                                                 45
<210> 283
<211> 18
<212> PRT
<213> Caenorhabditis elegans
<400> 283
Glu Ile Ala Ser Ala Ala Glu Thr Gly Trp Asp Phe Ser Thr Arg Trp
                 5
                                    10
1
Phe Ser
<210> 284
<211> 15
<212> PRT
<213> Caenorhabditis elegans
<400> 284
Ala Ser Ala Ala Glu Gly Trp Asp Phe Ser Thr Arg Trp Phe Ser
1
                                    10
                                                         15
<210> 285
<211> 18
```

```
<212> PRT
<213> Caenorhabditis elegans
<400> 285
Asp Leu Ala Ser Ala Ala Glu Ser Gly Trp Asp Phe Ser Thr Arg Trp
                 5
1
                                     10
Phe Ser
<210> 286
<211> 40
<212> PRT
<213> Caenorhabditis elegans
<400> 286
Lys Gln Phe Pro Tyr Tyr Gln Tyr Lys Ala Lys Leu Lys Val Pro Arg
1
                                    10
Pro Glu Ser Tyr Arg Glu Asp Ser Glu Leu Ala Glu His Leu Gln Thr
            20
                                25
Glu Ala Glu Lys Ile Gln Met Trp
<210> 287
<211> 18
<212> PRT
<213> Caenorhabditis elegans
<400> 287
Lys Phe Tyr Gln Tyr Lys Val Pro Arg Pro Glu Ser Tyr Arg Asp Leu
                                                         15
Ala Gln
<210> 288
<211> 40
<212> PRT
<213> Caenorhabditis elegans
<400> 288
Lys Ser Phe Lys Val Tyr Gln Tyr Lys Thr Ala Ser Asn Val Pro Arg
1
                                    10
                                                         15
Pro Glu Ser Tyr Arg Val Asp Thr Gln Asn Ser Ala Lys Leu Ala Asn
            20
                                25
Gly Ala Asp Gln Gln Phe Tyr
        35
<210> 289
<211> 21
<212> PRT
<213> Caenorhabditis elegans
<400> 289
Gln Thr Gly Phe Gly Trp Thr Asn Gly Val Ile Leu Asp Leu Leu Asp
1
                                    10
```

Lys Tyr Gly Asp Gln

```
<210> 290
<211> 14
<212> PRT
<213> Caenorhabditis elegans
<400> 290
Gln Gly Phe Gly Trp Asn Gly Ile Leu Asp Leu Leu Tyr Asp
                                    10
<210> 291
<211> 21
<212> PRT
<213> Caenorhabditis elegans
<400> 291
Gln Asp Gly Phe Gly Trp Ser Asn Gly Ala Ile Leu Asp Leu Leu
1
                                    10
Thr Tyr Asn Asp Arg
            20
<210> 292
<211> 27
<212> PRT
<213> Caenorhabditis elegans
<400> 292
Tyr Gly Asp Gln Phe Ala Ser Ser Ser Thr Ala Ser Lys Phe Ser Phe
1
                                    10
Ser Leu Ser Asn Ile Thr Phe Val Val Phe Ile
            20
<210> 293
<211> 11
<212> PRT
<213> Caenorhabditis elegans
<400> 293
Tyr Phe Ala Ser Ser Ser Ala Ser Phe Ser Phe
                 5
<210> 294
<211> 26
<212> PRT
<213> Caenorhabditis elegans
<400> 294
Tyr Asn Pro Phe Ala Ser Ser Ser Asp Ala Ser Ser Cys Pro Phe Ser
1
                                    10
Thr Asn Ser Val Ile Phe Ser Ile Leu Val
            20
                                25
```

```
<210> 295
<211> 9
<212> PRT
<213> Caenorhabditis elegans
<400> 295
Gly Gly Gly Glu Tyr Glu Val Gln
                 5
<210> 296
<211> 7
<212> PRT
<213> Caenorhabditis elegans
<400> 296
Gly Gly Glu Tyr Val Gln
<210> 297
<211> 9
<212> PRT
<213> Caenorhabditis elegans
<400> 297
Gly Ser Gly Gly Glu Tyr Asp Val Gln
1
                 5
<210> 298
<211> 14
<212> PRT
<213> Caenorhabditis elegans
<400> 298
Asn Gln Tyr Tyr Val Ser Asn Ala Val Pro Leu Tyr Ala Lys
                 5
<210> 299
<211> 7
<212> PRT
<213> Caenorhabditis elegans
<400> 299
Asn Tyr Tyr Val Leu Tyr Lys
1
<210> 300
<211> 14
<212> PRT
<213> Caenorhabditis elegans
<400> 300
Asn His Tyr Tyr Ile Ile Gln Met Val Ser Leu Tyr Thr Lys
                 5
```

```
<210> 301
<211> 30
<212> PRT
<213> Caenorhabditis elegans
<400> 301
Asp Gln Phe Ala Ser Ser Ser Thr Ala Ser Lys Phe Ser Phe Ser Leu
                5
                                     10
Ser Asn Ile Thr Phe Val Val Phe Ile Leu Tyr Ile Phe Ser
            20
                                25
                                                     30
<210> 302
<211> 11
<212> PRT
<213> Caenorhabditis elegans
<400> 302
Asp Gln Phe Ser Ser Lys Phe Ser Phe Phe Ser
                 5
<210> 303
<211> 30
<212> PRT
<213> Caenorhabditis elegans
<400> 303
Asp Gln Phe Val Ile Ser Phe Ile Cys Ser Lys Phe Ser Ser Lys Asn
1
                5
                                     10
Lys Lys Leu Tyr Phe Cys Pro Ser His Phe Ser Leu Phe Ser
            20
                                25
                                                     30
<210> 304
<211> 9
<212> PRT
<213> Caenorhabditis elegans
<220>
<221> VARIANT
<222> (1)...(9)
<223> Xaa = Any Amino Acid
<400> 304
Gly Trp Asp Xaa Xaa Ile Ala Pro Lys
<210> 305
<211> 62
<212> PRT
<213> Mus musculus
<400> 305
Leu Cys Lys Glu Gly Ile Ser Asp Gly Ala Thr Met Lys Thr Phe Cys
Gly Thr Pro Glu Tyr Leu Ala Pro Glu Val Leu Glu Asp Asn Asp Tyr
```

30

25

```
Gly Arg Ala Val Asp Trp Trp Gly Leu Gly Val Val Met Tyr Glu Met
                            40
Met Cys Gly Arg Leu Pro Phe Tyr Asn Gln Asp His Glu Arg
                        55
<210> 306
<211> 9
<212> PRT
<213> Caenorhabditis elegans
<400> 306
Gln Ala Leu Thr Gln Met Asn Pro Lys
                 5
<210> 307
<211> 11
<212> PRT
<213> Caenorhabditis elegans
<400> 307
Gln Ala Leu Thr Gln Cys Val Asp Ser Met Arg
                 5
<210> 308
<211> 248
<212> PRT
<213> Homo sapiens
<400> 308
Ile Phe Arg Thr Ala Val Ser Ser Asn Arg Cys Arg Thr Glu Tyr Gln
                5
                                    10
Asn Ile Asp Leu Asp Cys Ala Tyr Ile Thr Asp Arg Ile Ile Ala Ile
                                25
Gly Tyr Pro Ala Thr Gly Ile Glu Ala Asn Phe Arg Asn Ser Lys Val
                            40
                                                 45
Gln Thr Gln Gln Phe Leu Thr Arg Arg His Gly Lys Gly Asn Val Lys
                        55
                                            60
Val Phe Asn Leu Arg Gly Gly Tyr Tyr Tyr Asp Ala Asp Asn Phe Asp
                    70
                                        75
Gly Asn Val Ile Cys Phe Asp Met Thr Asp His His Pro Pro Ser Leu
                85
                                    90
Glu Leu Met Ala Pro Phe Cys Arg Glu Ala Lys Glu Trp Leu Glu Ala
                                105
Asp Asp Lys His Val Ile Ala Val His Cys Lys Ala Gly Lys Gly Arg
        115
                            120
Thr Gly Val Met Ile Cys Ala Leu Leu Ile Tyr Ile Asn Phe Tyr Pro
                        135
                                            140
Ser Pro Arg Gln Ile Leu Asp Tyr Tyr Ser Ile Ile Thr Arg Lys Asn
Asn Lys Gly Val Thr Ile Pro Ser Gln Arg Arg Tyr Ile Tyr Tyr Tyr
                                    170
His Lys Leu Arg Glu Arg Glu Leu Asn Tyr Leu Pro Leu Arg Met Gln
                                185
Leu Ile Gly Val Tyr Val Glu Arg Pro Pro Lys Thr Trp Gly Gly Gly
                            200
```

Ser Lys Ile Lys Val Glu Val Gly Asn Gly Ser Thr Ile Leu Phe Lys

```
215
Pro Asp Pro Leu Ile Ile Ser Lys Ser Asn His Gln Arg Glu Arg Ala
                   230
                                        235
Thr Trp Leu Asn Asn Cys Asp Thr
<210> 309
<211> 249
<212> PRT
<213> Caenorhabditis elegans
<400> 309
Ile Ile Lys Glu Ile Val Ser Arg Asn Lys Arg Arg Tyr Gln Glu Asp
                                    10
Gly Phe Asp Leu Asp Leu Thr Tyr Ile Tyr Pro Asn Ile Ile Ala Met
                                25
Gly Phe Pro Ala Glu Arg Leu Glu Gly Val Tyr Arg Asn Asn Ile Asp
                            40
Asp Val Val Arg Phe Leu Asp Ser Lys His Lys Asn His Tyr Lys Ile
                        55
Tyr Asn Leu Cys Ala Glu Arg His Tyr Asp Thr Ala Lys Phe Asn Cys
                   70
                                        75
Arg Val Ala Gln Tyr Pro Phe Glu Asp His Asn Pro Pro Gln Leu Glu
               85
                                    90
Leu Ile Lys Pro Phe Cys Glu Asp Leu Asp Gln Trp Leu Ser Glu Asp
           100
                                105
                                                    110
Asp Asn His Val Ala Ala Ile His Cys Lys Ala Gly Lys Gly Arg Thr
       115
                           120
                                                125
Gly Val Met Ile Cys Ala Tyr Leu Leu His Arg Gly Lys Phe Leu Lys
                        135
                                            140
Ala Gln Glu Ala Leu Asp Phe Tyr Gly Glu Val Arg Thr Arg Asp Lys
                   150
                                        155
Lys Gly Val Thr Ile Pro Ser Gln Arg Arg Tyr Val Tyr Tyr Ser
               165
                                    170
                                                        175
Tyr Leu Leu Lys Asn His Leu Asp Tyr Arg Pro Val Ala Leu Leu Phe
           180
                                185
His Lys Met Met Phe Glu Thr Ile Pro Met Phe Ser Gly Gly Thr Cys
                           200
                                                205
Asn Pro Gln Phe Val Val Cys Gln Leu Lys Val Lys Ile Tyr Ser Ser
                                           220
                        215
Asn Ser Gly Pro Thr Arg Arg Glu Asp Lys Phe Asn Tyr Phe Glu Phe
                   230
                                        235
Pro Gln Pro Leu Pro Val Cys Gly Asp
                245
<210> 310
<211> 962
<212> PRT
<213> Caenorhabditis elegans
<400> 310
Met Val Thr Pro Pro Pro Asp Val Pro Ser Thr Ser Thr Arg Ser Met
                                    10
Ala Arg Asp Leu Gln Glu Asn Pro Asn Arg Gln Pro Gly Glu Pro Arg
                               25
Val Ser Glu Pro Tyr His Asn Ser Ile Val Glu Arg Ile Arg His Ile
```

Phe Arg Thr Ala Val Ser Ser Asn Arg Cys Arg Thr Glu Tyr Gln Asn Ile Asp Leu Asp Cys Ala Tyr Ile Thr Asp Arg Ile Ile Ala Ile Gly Tyr Pro Ala Thr Gly Ile Glu Ala Asn Phe Arg Asn Ser Lys Val Gln Thr Gln Gln Phe Leu Thr Arg Arg His Gly Lys Gly Asn Val Lys Val Phe Asn Leu Arg Gly Gly Tyr Tyr Tyr Asp Ala Asp Asn Phe Asp Gly Asn Val Ile Cys Phe Asp Met Thr Asp His His Pro Pro Ser Leu Glu Leu Met Ala Pro Phe Cys Arg Glu Ala Lys Glu Trp Leu Glu Ala Asp Asp Lys His Val Ile Ala Val His Cys Lys Ala Gly Lys Gly Arg Thr Gly Val Met Ile Cys Ala Leu Leu Ile Tyr Ile Asn Phe Tyr Pro Ser Pro Arg Gln Ile Leu Asp Tyr Tyr Ser Ile Ile Arg Thr Lys Asn Asn Lys Gly Val Thr Ile Pro Ser Gln Arg Arg Tyr Ile Tyr Tyr His Lys Leu Arg Glu Arg Glu Leu Asn Tyr Leu Pro Leu Arg Met Gln Leu Ile Gly Val Tyr Val Glu Arg Pro Pro Lys Thr Trp Gly Gly Gly Ser Lys Ile Lys Val Glu Val Gly Asn Gly Ser Thr Ile Leu Phe Lys Pro Asp Pro Leu Ile Ile Ser Lys Ser Asn His Gln Arg Glu Arg Ala Thr Trp Leu Asn Asn Cys Asp Thr Pro Asn Glu Phe Asp Thr Gly Glu Gln Lys Tyr His Gly Phe Val Ser Lys Arg Ala Tyr Cys Phe Met Val Pro Glu Asp Ala Pro Val Phe Val Glu Gly Asp Val Arg Ile Asp Ile Arg Glu Ile Gly Phe Leu Lys Lys Phe Ser Asp Gly Lys Ile Gly His Val Trp Phe Asn Thr Met Phe Ala Cys Asp Gly Gly Leu Asn Gly Gly His Phe Glu Tyr Val Asp Lys Thr Gln Pro Tyr Ile Gly Asp Asp Thr Ser Ile Gly Arg Lys Asn Gly Met Arg Arg Asn Glu Thr Pro Met Arg Lys Ile Asp Pro Glu Thr Gly Asn Glu Phe Glu Ser Pro Trp Gln Ile Val Asn Pro Pro Gly Leu Glu Lys His Ile Thr Glu Glu Gln Ala Met Glu Asn Tyr Thr Asn Tyr Gly Met Ile Pro Pro Arg Tyr Thr Ile Ser Lys Ile Leu His Glu Lys His Glu Lys Gly Ile Val Lys Asp Asp Tyr Asn Asp Arg Lys Leu Pro Met Gly Asp Lys Ser Tyr Thr Glu Ser Gly Lys Ser Gly Asp Ile Arg Gly Val Gly Gly Pro Phe Glu Ile Pro Tyr Lys Ala Glu Glu His Val Leu Thr Phe Pro Val Tyr Glu Met Asp Arg Ala Leu Lys Ser Lys Asp Leu Asn Asn Gly Met Lys Leu His Val Val Leu

```
515
                            520
Arg Cys Val Asp Thr Arg Asp Ser Lys Met Met Glu Lys Ser Glu Val
    530
                        535
                                             540
Phe Gly Asn Leu Ala Phe His Asn Glu Ser Thr Arg Arg Leu Gln Ala
                    550
                                         555
Leu Thr Gln Met Asn Pro Lys Trp Arg Pro Glu Pro Cys Ala Phe Gly
                565
                                    570
Ser Lys Gly Ala Glu Met His Tyr Pro Pro Ser Val Arg Tyr Ser Ser
            580
                                585
                                                     590
Asn Asp Gly Lys Tyr Asn Gly Ala Cys Ser Glu Asn Leu Val Ser Asp
        595
                            600
Phe Phe Glu His Arg Asn Ile Ala Val Leu Asn Arg Tyr Cys Arg Tyr
                        615
Phe Tyr Lys Gln Arg Ser Thr Ser Arg Ser Arg Tyr Pro Arg Lys Phe
                    630
                                         635
Arg Tyr Cys Pro Leu Ile Lys Lys His Phe Tyr Ile Pro Ala Asp Thr
                645
                                    650
Asp Asp Val Asp Glu Asn Gly Gln Pro Phe Phe His Ser Pro Glu His
            660
                                665
Tyr Ile Lys Glu Gln Glu Lys Ile Asp Ala Glu Lys Ala Ala Lys Gly
        675
                            680
Ile Glu Asn Thr Gly Pro Ser Thr Ser Gly Ser Ser Ala Pro Gly Thr
                        695
Ile Lys Lys Thr Glu Ala Ser Gln Ser Asp Lys Val Lys Pro Ala Thr
                    710
                                        715
Glu Asp Glu Leu Pro Pro Ala Arg Leu Pro Asp Asn Val Arg Arg Phe
                725
                                    730
Pro Val Val Gly Val Asp Phe Glu Asn Pro Glu Glu Glu Ser Cys Glu
            740
                                745
His Lys Thr Val Glu Ser Ile Ala Gly Phe Glu Pro Leu Glu His Leu
                            760
Phe His Glu Ser Tyr His Pro Asn Thr Ala Gly Asn Met Leu Arg Gln
                        775
Asp Tyr His Thr Asp Ser Glu Val Lys Ile Ala Glu Gln Glu Ala Lys
                    790
                                        795
Ala Phe Val Asp Gln Leu Leu Asn Gly Gln Gly Val Leu Gln Glu Phe
                805
                                    810
Met Lys Gln Phe Lys Val Pro Ser Asp Asn Ser Phe Ala Asp Tyr Val
            820
                                825
Thr Gly Gln Ala Glu Val Phe Lys Ala Gln Ile Ala Leu Leu Glu Gln
                            840
Ser Glu Asp Phe Gln Arg Val Gln Ala Asn Ala Glu Glu Val Asp Leu
                        855
Glu His Thr Leu Gly Glu Ala Phe Glu Arg Phe Gly His Val Val Glu
                    870
                                        875
Glu Ser Asn Gly Ser Ser Lys Asn Pro Lys Ala Leu Lys Thr Arg Glu
                885
                                    890
Gln Met Val Lys Glu Thr Gly Lys Asp Thr Gln Lys Thr Arg Asn His
            900
                                905
Val Leu Leu His Leu Glu Ala Asn His Arg Val Gln Ile Glu Arg Arg
                            920
Glu Thr Cys Pro Glu Leu His Pro Glu Asp Lys Ile Pro Arg Ile Ala
                        935
                                            940
His Phe Ser Glu Asn Ser Phe Ser Asp Ser Asn Phe Asp Gln Ala Ile
945
                    950
Tyr Leu
```

```
<210> 311
<211> 3304
<212> DNA
<213> Caenorhabditis elegans
```

<400> 311

ttccaggtac atctactaac ccccaatggt tactcctcct ccagatgtgc caagcacatc 60 120 gaccaggtcg atggctcgtg accttcaaga gaatccaaac cgacaacctg gtgaaccacg tgtgtctgaa ccgtatcaca attcaatcgt cgagcggatt cgccatattt ttcggacggc 180 240 tgtatcttcc aatcgttgtc gcaccgagta ccaaaatatc gacctagatt gtgcatatat cacagaccga atcatagcta tcggttatcc agcaacagga atcgaagcga atttccgtaa 300 ctcaaaagtt caaactcaac aatttctgac caggcggcac ggaaagggca acgtgaaggt 360 gtttaacctg cgcggtggat actactacga tgcggataac ttcgatggaa atgttatttq 420 cttcgatatg actgatcatc atccgccgag tctcgaatta atggctccgt tttgcagaga 480 ggctaaggaa tggcttgaag cagacgataa acatgtaata gctgtacact gtaaagctgg 540 aaaaggccgt accggagtga tgatatgtgc tcttctcatc tacatcaact tctatccgag 600 cccacgacaa attctcgact actactcaat aattcgtaca aaaaacaaca aaggtgtcac 660 aattccatca caacgacgct acatttacta ctaccataag cttcgtgaac gtgagctcaa 720 780 ctatttacca ttgagaatgc agttgattgg tgtctacgtg gaacggcctc caaagacatg gggtggtggt tcaaagataa aagtggaggt tggaaatggc tcgacaattt tatttaagcc 840 ggatcctctc ataatctcca aatcaaatca tcagcgagag cgtgcgacgt ggctgaacaa 900 ctgtgatacg cctaacgaat tcgacaccgg agagcaaaaa tatcatggat ttgtttccaa 960 gagagcatac tgttttatgg tgccagaaga tgctccagta tttgtcgaag gagatgttcg 1020 tatagacatt cgcgaaatcg gatttctcaa aaagttttcg gacgggaaga ttggtcatgt 1080 1140 ttggttcaat acaatgttcg catgtgatgg aggactcaac ggtggacatt tcgagtacgt agacaaaact cagccgtaca tcggagacga tacatcaatc ggacggaaaa atggaatgcg 1200 aagaaatgaa acgccgatgc gaaaaattga tccagaaact ggaaatgaat ttgagtctcc 1260 gtggcaaata gtgaatcctc ctggactgga aaaacatatt acggaggaac aagcaatgga 1320 aaattatacc aattatggca tgattcctcc tcgatacacg atcagcaaga ttcttcacga 1380 1440 aaagcatgaa aaaggtatcg tcaaggatga ctataatgat cgtaagctgc caatgggaga 1500 caaatcatac acggaatcag gaaaaagtgg agatattcga ggagtcggtg gtccatttga gataccatat aaagctgagg aacatgttct cacatttcca gtttatgaaa tggatcgagc 1560 attgaagagt aaagatetta acaacggaat gaaacttcac gttgttette gttgtgtaga 1620 tactcgtgat tcaaaaatga tggaaaagag cgaagtgttc ggcaatctgg cattccataa 1680 tgaatcgaca cggaggcttc aagcgttgac tcaaatgaat ccaaaatggc gacctgaacc 1740 gtgtgcgttc ggatccaaag gtgctgaaat gcattaccct ccgtcggttc gatattcaag 1800 caatgatgga aagtataatg gagcctgcag tgagaacctt gttagcgatt ttttcgagca 1860 cagaaatatt gccgttctta atcgatattg ccgatatttc tacaagcaac gcagtacatc 1920 tcgaagccgt tatccaagaa aattcagata ctgtcctctg atcaagaaac atttctacat 1980 tocagotgat accgatgatg ttgatgaaaa tgggcaaccg ttcttccact caccagagca 2040 2100 ttacattaaa gaacaggaaa aaatagacgc agagaaagca gctaaaggaa ttgaaaatac tggacccagt acttcaggat caagtgctcc cggaactatc aagaaaacgg aagcttcaca 2160 2220 atccgacaag gtgaagccgg caactgaaga cgaacttcct cctgcgaggc taccggataa tgtgcgaaga tttccagtcg tcggcgttga tttcgaaaat ccggaagaag aatcgtgtga 2280 acacaaaacc gtagagtcaa tagctggttt tgaaccactc gaacatctat tccatgaatc 2340 ataccatcca aatacggccg gtaacatgct gcgtcaggat tatcacactg attcggaagt 2400 gaaaatagct gaacaagagg caaaagcctt cgttgaccag ttgcttaatg gacaaggtgt 2460 2520 attacaagag tttatgaagc aattcaaagt accatcggac aattcctttg ctgattatgt 2580 aaccggacag gccgaagttt ttaaagcaca gattgcgtta ctggagcagt cggaggattt 2640 tcaacgagtt caagcgaatg cagaggaagt cgatcttgaa cacactcttg gtgaagcgtt tgagcgattc gggcacgttg tagaagaatc gaatggttct tctaaaaaatc caaaagccct 2700 gaaaactcga gaacaaatgg tgaaagaaac tggcaaagac actcagaaga cccgcaatca 2760 tgtgcttcta catttggaag ctaatcatcg tgtgcaaatc gagcgtcgtg aaacgtgccc 2820 ggagctacat ccagaggata aaatcccaag aattgctcat ttttccgaaa acagcttctc 2880 ggattcgaat tttgatcaag ctatttattt gtaaacctaa aacaaaactt ttagaagatt 2940 ttcttcttac tgaccctcca attttcagat aatttcaatg ttttaagttt tctcttcaaa 3000 gtatcattca ctttctgtat agtgttttgt tttttaacaa actattgttc gattattttg 3060 tatattcata ttatagctct caacttcccg attttccacg tatatatgta tattttgccg 3120 ggtgaaaaat agcaattccc tatgaatgta tccccttcca tctgttttct tactcagaaa 3180

```
ttgtaattca cattgcgggt catcactaat cctatgggct ttaacacaat tctcccataa
                                                                      3240
attaattqta cttaccaatt ttttqtttaa ttatttaqat ttqtaacatt gaaattqqtq
                                                                      3300
                                                                      3304
ataa
<210> 312
<211> 1642
<212> DNA
<213> Caenorhabditis elegans
<400> 312
tttaattacc caagtttgag gtagcattgc tctcttcaat catatggatt cgttgtttca
                                                                        60
gatggcatcc gcaatgaagt ttcaatacta ctcgaagaaa gctgctggaa agacaatgtc
                                                                       120
taatagtgtc tccatgtcca gtgacaatcg catggaggat tttaaacgtc gttttcgtcg
                                                                       180
aagtggatcg ttaggaattc catttgtccc agaagaagat gttaaacaac tcttcacacc
                                                                       240
aactcgtact gttcgtcgag aagcatctat tcgcgaaggg gatgaggaag aaggagtaca
                                                                       300
                                                                       360
aattctcaca ataattgtca agtcaagtcg tgtttcggag gatatctcaa aaatgattgc
aaacctccct gatcacactc gtatcaaaca tttggagact cgtgacagtc aagatggaag
                                                                       420
                                                                       480
ttccaaaact atggatgttc ttctagagat tgagctcttt cattatggaa aacaagaagc
aatggatctt atgagactta atgggcttga tgttcatgag gtgtcatcga ctattcgtcc
                                                                       540
                                                                       600
aactgcaata aaagagcaat atacagagcc tggatctgat gatgcgacaa ccggttctga
                                                                       660
atggtttcca aaaagtattt atgatttgga tatttgtgca aaaagagtga ttatgtatgg
agcagggctg gacgctgatc atcctggttt caaagatacc gagtatcgtc aacgtcgaat
                                                                       720
gatgtttgct gaactggcgc tcaattacaa acacggtgag ccaattccgc gaaccgaata
                                                                       780
                                                                       840
tacatcatcc gaacggaaaa cttggggaat tatatataga aaattgagag aattgcacaa
aaagcacgca tgcaagcagt ttcttgataa ctttgagcta ctggagagac attgtggata
                                                                       900
                                                                       960
ctcggaaaat aatattccgc aactagaaga tatctgcaag tttttgaaag caaaaactgg
attecgtgtt egeceagteg eeggataett ateagetegt gatttettgg eaggtettge
                                                                      1020
                                                                      1080
atatcgtgtc ttcttctgca ctcaatacgt tcgccatcat gccgatccat tttacactcc
                                                                      1140
agaaccagac accgttcacg agctcatggg tcacatggct ctattcgctg atccagattt
tgctcagttt tctcaagaga ttggattagc ttctcttgga gcatcagagg aagatttgaa
                                                                      1200
gaagettgca acactetact tettttecat tgaatttggt etetegtetg atgaegetge
                                                                      1260
cgattctcca gtaaaagaaa atggatcaaa tcatgaaaga tttaaaagtat acggagcagg
                                                                      1320
acttctgagc agtgctggcg agttgcaaca tgccgttgag ggtagtgcaa ccattattcg
                                                                      1380
ttttgatccg gatcgtgttg ttgagcaaga atgtctcatt actactttcc agtcagcgta
                                                                      1440
tttctatact agaaattttg aagaggccca gcagaaactc agaatgttca ccaacaacat
                                                                      1500
qaaacqtccc ttcattqttc qttacaaccc atacacagaa agcqtcgaag ttctcaacaa
                                                                      1560
ctcccqttcc attatqttqq caqtqaactc tctccqctca gacatcaacc tgctcqccgg
                                                                      1620
                                                                      1642
agctctccac tacatcctgt ag
<210> 313
<211> 532
<212> PRT
<213> Caenorhabditis elegans
Met Asp Ser Leu Phe Gln Met Ala Ser Ala Met Lys Phe Gln Tyr Tyr
1
                                    10
Ser Lys Lys Ala Ala Gly Lys Thr Met Ser Asn Ser Val Ser Met Ser
                                25
Ser Asp Asn Arg Met Glu Asp Phe Lys Arg Arg Phe Arg Arg Ser Gly
                            40
Ser Leu Gly Ile Pro Phe Val Pro Glu Glu Asp Val Lys Gln Leu Phe
Thr Pro Thr Arg Thr Val Arg Arg Glu Ala Ser Ile Arg Glu Gly Asp
Glu Glu Gly Val Gln Ile Leu Thr Ile Ile Val Lys Ser Ser Arg
                                    90
Val Ser Glu Asp Ile Ser Lys Met Ile Ala Asn Leu Pro Asp His Thr
```

105

Arg Ile Lys His Leu Glu Thr Arg Asp Ser Gln Asp Gly Ser Ser Lys Thr Met Asp Val Leu Leu Glu Ile Glu Leu Phe His Tyr Gly Lys Gln Glu Ala Met Asp Leu Met Arg Leu Asn Gly Leu Asp Val His Glu Val Ser Ser Thr Ile Arg Pro Thr Ala Ile Lys Glu Gln Tyr Thr Glu Pro Gly Ser Asp Asp Ala Thr Thr Gly Ser Glu Trp Phe Pro Lys Ser Ile Tyr Asp Leu Asp Ile Cys Ala Lys Arg Val Ile Met Tyr Gly Ala Gly Leu Asp Ala Asp His Pro Gly Phe Lys Asp Thr Glu Tyr Arg Gln Arg Arg Met Met Phe Ala Glu Leu Ala Leu Asn Tyr Lys His Gly Glu Pro Ile Pro Arg Thr Glu Tyr Thr Ser Ser Glu Arg Lys Thr Trp Gly Ile Ile Tyr Arg Lys Leu Arg Glu Leu His Lys Lys His Ala Cys Lys Gln Phe Leu Asp Asn Phe Glu Leu Leu Glu Arg His Cys Gly Tyr Ser Glu Asn Asn Ile Pro Gln Leu Glu Asp Ile Cys Lys Phe Leu Lys Ala Lys Thr Gly Phe Arg Val Arg Pro Val Ala Gly Tyr Leu Ser Ala Arg Asp Phe Leu Ala Gly Leu Ala Tyr Arg Val Phe Phe Cys Thr Gln Tyr Val Arg His His Ala Asp Pro Phe Tyr Thr Pro Glu Pro Asp Thr Val His Glu Leu Met Gly His Met Ala Leu Phe Ala Asp Pro Asp Phe Ala Gln Phe Ser Gln Glu Ile Gly Leu Ala Ser Leu Gly Ala Ser Glu Glu Asp Leu Lys Lys Leu Ala Thr Leu Tyr Phe Phe Ser Ile Glu Phe Gly Leu Ser Ser Asp Asp Ala Ala Asp Ser Pro Val Lys Glu Asn Gly Ser Asn His Glu Arg Phe Lys Val Tyr Gly Ala Gly Leu Leu Ser Ser Ala Gly Glu Leu Gln His Ala Val Glu Gly Ser Ala Thr Ile Ile Arg Phe Asp Pro Asp Arg Val Val Glu Glu Cys Leu Ile Thr Thr Phe Gln Ser Ala Tyr Phe Tyr Thr Arg Asn Phe Glu Glu Ala Gln Gln Lys Leu Arg Met Phe Thr Asn Asn Met Lys Arg Pro Phe Ile Val Arg Tyr Asn Pro Tyr Thr Glu Ser Val Glu Val Leu Asn Asn Ser Arg Ser Ile Met Leu Ala Val Asn Ser Leu Arg Ser Asp Ile Asn Leu Leu Ala Gly Ala Leu His Tyr Ile Leu

<210> 314

<211> 817

<212> DNA

<213> Caenorhabditis elegans

```
<400> 314
                                                                        60
attacccaag tttgaggtag cattgctctc ttcaatcata tggattcgtt gtttcagatg
                                                                       120
gcatccgcaa tgaagtttca atactactcg aagaaagctg ctggaaagac aatgtctaat
                                                                       180
agtgtcaaaa actggattcc gtgttcgccc agtcgccgga tacttatcag ctcgtgattt
                                                                       240
cttggcaggt cttgcatatc gtgtcttctt ctgcactcaa tacgttcgcc atcatgccga
tocattttac actocagaac cagacaccgt toacgagete atgggtcaca tggctctatt
                                                                       300
                                                                       360
cgctgatcca gattttgctc agttttctca agagattgga ttagcttctc ttggagcatc
                                                                       420
agaggaagat ttgaagaagc ttgcaacact ctacttcttt tccattgaat ttggtctctc
                                                                       480
gtctgatgac gctgccgatt ctccagtaaa agaaaatgga tcaaatcatg aaagatttaa
agtatacgga gcaggacttc tgagcagtgc tggcgagttg caacatgccg ttgagggtag
                                                                       540
tgcaaccatt attcgttttg atccggatcg tgttgttgag caagaatgtc tcattactac
                                                                       600
tttccagtca gcgtatttct atactagaaa ttttgaagag gcccagcaga aactcagaat
                                                                       660
gttcaccaac aacatgaaac gtcccttcat tgttcgttac aacccataca cagaaagcgt
                                                                       720
                                                                       780
cgaagttctc aacaactccc gttccattat gttggcagtg aactctctcc gctcagacat
                                                                       817
caacctgctc gccggagctc tccactacat cctgtag
<210> 315
<211> 45
<212> PRT
<213> Caenorhabditis elegans
<400> 315
Met Asp Ser Leu Phe Gln Met Ala Ser Ala Met Lys Phe Gln Tyr Tyr
1
                 5
                                    10
                                                         15
Ser Lys Lys Ala Ala Gly Lys Thr Met Ser Asn Ser Val Lys Asn Trp
                                25
                                                     30
Ile Pro Cys Ser Pro Ser Arg Arg Ile Leu Ile Ser Ser
                            40
<210> 316
<211> 466
<212> DNA
<213> Caenorhabditis elegans
<400> 316
                                                                        60
atteggeatg ageatggage ttegagteet agagaacaca aaaegtteee ggeggaacet
                                                                       120
gggtctggac tgcgacgaga ctcaagcgag tcccgctgct gccgatatcc cctcacagtg
                                                                       180
gactttgagg ctttcggctg ggactggatc atcgcaccta agcgctacaa ggccaactac
tgctccggcc agtgggagta catgttcatg caaaaatatc cgcataccca tttggtgcag
                                                                       240
                                                                       300
caggccaatc caagaggtta tgctgggccc tgttgtaccc ccaccaagat gtccccaatc
                                                                       360
aacatgctct acttcaatga caagcagcag attatctacg gcaagatccc tggcatggtg
                                                                       420
qtqqatcqct qtqqctqctc ttaaqqtqqq qqataqaqqa tqcctccccc acaqaccqta
                                                                       466
ccccaagacc catagccctg cccaatccac cgcctgatcc aaacat
<210> 317
<211> 128
<212> PRT
<213> Caenorhabditis elegans
<400> 317
Ile Arg His Glu His Gly Ala Ser Ser Pro Arg Glu His Lys Thr Phe
                                                         15
                                     10
Pro Ala Glu Pro Gly Ser Gly Leu Arg Arg Asp Ser Ser Glu Ser Arg
                                25
Cys Cys Arg Tyr Pro Leu Thr Val Asp Phe Glu Ala Phe Gly Trp Asp
                            40
```

```
Trp Ile Ile Ala Pro Lys Arg Tyr Lys Ala Asn Tyr Cys Ser Gly Gln
    50
                        55
Trp Glu Tyr Met Phe Met Gln Lys Tyr Pro His Thr His Leu Val Gln
                                         75
Gln Ala Asn Pro Arg Gly Tyr Ala Gly Pro Cys Cys Thr Pro Thr Lys
Met Ser Pro Ile Asn Met Leu Tyr Phe Asn Asp Lys Gln Gln Ile Ile
                                 105
                                                     110
Tyr Gly Lys Ile Pro Leu Ala Met Val Val Asp Arg Cys Gly Cys Ser
                            120
<210> 318
<211> 9
<212> DNA
<213> Homo sapiens
<400> 318
                                                                          9
caaaactaa
<210> 319
<211> 20
<212> DNA
<213> Caenorhabditis elegans
<400> 319
                                                                         20
ccactatggc cgagatttcc
<210> 320
<211> 44
<212> DNA
<213> Caenorhabditis elegans
<400> 320
                                                                         44
ccagtgaaaa gttcttctcc tttcttcctc ttctcgaatt cgga
<210> 321
<211> 21
<212> DNA
<213> Caenorhabditis elegans
<400> 321
cttcctcttc tcgaattcgg c
                                                                         21
<210> 322
<211> 8
<212> PRT
<213> Caenorhabditis elegans
<400> 322
Gly Arg Lys Gly Phe Pro His Val
                 5
<210> 323
<211> 7
<212> PRT
<213> Caenorhabditis elegans
```

```
<220>
<221> VARIANT
<222> (1) ... (7)
<223> Xaa = Any Amino Acid
<400> 323
Arg Xaa Xaa Ile Xaa Xaa Gly
<210> 324
<211> 7
<212> PRT
<213> Caenorhabditis elegans or Homo sapiens
<400> 324
Cys Gly Cys Cys Cys Cys
<210> 325
<211> 79
<212> PRT
<213> Homo sapiens or Caenorhabditis elegans
<400> 325
Val Leu Asp Asp Tyr Gly Arg Val Asp Trp Trp Gly Gly Val Val Met
                                    10
Tyr Glu Met Met Cys Gly Arg Leu Pro Phe Tyr Asp His Lys Leu Phe
            20
                                25
                                                    30
Glu Leu Ile Arg Phe Pro Leu Glu Ala Leu Leu Gly Leu Leu Lys Asp
                            40
Pro Thr Gln Arg Leu Gly Gly Glu Asp Ala Glu Ile Phe Trp
                        55
Tyr Lys Pro Pro Lys Pro Val Ser Glu Thr Asp Thr Tyr Phe Asp
<210> 326
<211> 48
<212> PRT
<213> Homo sapiens or Caenorhabditis elegans
<400> 326
Thr Met Phe Leu Lys Leu Gly Lys Gly Thr Phe Gly Lys Val Ile Leu
                                                        15
                                    10
Lys Glu Lys Thr Tyr Ala Lys Ile Leu Lys Lys Val Ile Ala Glu Val
            20
                                25
Ala His Thr Leu Thr Glu Asn Arg Val Leu Gln His Pro Phe Leu Thr
                            40
<210> 327
<211> 27
<212> DNA
<213> Caenorhabditis elegans
<400> 327
caagcgttga ctcaaatgaa tccaaaa
```

```
<210> 328
<211> 55
<212> DNA
```

<213> Caenorhabditis elegans

<400> 328

caagcgttga ctcaatgcgt tgactcaatg cgttgactcg ttgacgaatc caaaa